



SEQUENCE LISTING

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<1> Luparello, James R
Koel, Cornelius F
Takashima, Hiroshi

<120> Defects in Periaxin Associated with Myelinopathies

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<140> US 10/021,955

<141> 2001-12-13

<150> US 60/255,217

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SEQUENCE LISTING

<110> Lupski, James R
Boerkoel, Cornelius F
Takashima, Hiroshi

<120> Defects in Periaxin Associated with Myelinopathies

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| gctgctgctc | ctcccccaag | gaaggccaag | gcagaagctg | aggcagccac | aggagctggg | 1560 |
| ttcacagccc | ctcagataga | gctagtgtgg | cctcggctgc | ctagtgccga | ggtgggtgtc | 1620 |
| cctcaggtct | cagttcccaa | ggggacccca | tcaacagagg | cagccagcgg | ctttgccctt | 1680 |
| cacctgccaa | cccttgggct | aggtgccccca | gctgcaccgg | ctgtggagcc | cccagccacg | 1740 |
| ggaatccagg | ttccacaagt | ggaactcccc | accctgcct | ctctaccac | gcttcccaca | 1800 |
| cttccatgcc | tggacaccca | ggaaggagct | gcagtggtaa | aagtccctac | cctggatgtg | 1860 |
| gcagctccgt | ctatgggggt | ggacctggct | ttgccgggtg | cagaggtgga | ggcccaggga | 1920 |
| gaggttcctg | aagtggccct | caagatgccc | cggctcagtt | tccccgttt | tgggattcgg | 1980 |
| gggaaggaag | ccactgaagc | caaagtagtc | aagggcagcc | ctgaggccaa | agcaaagggt | 2040 |
| cccagacttc | gaatgcccac | ctttgggctt | tctctcctgg | aaccccggcc | ctctggccct | 2100 |
| gaagctgttg | ctgagagcaa | gctgaagcta | cccaccctca | agatgccctc | tttcggcatt | 2160 |
| ggtgtggctg | ggcctgaagt | caaggcaccc | acggggcccg | aagtaaagct | ccctaagggt | 2220 |
| cctgaggtca | aactcccga | agtgcctgag | gcagccattc | cagatgtgca | actccctgag | 2280 |
| gtacagctgc | ccaaaatgtc | agacatgaaa | cttccaaaga | tccctgagat | ggttgtagcc | 2340 |
| gacgttcgtc | ttccggaagt | gcagctgccc | aaagtccctg | agatgaaagt | cccagagatg | 2400 |
| aagctcccga | agtggcccga | gatggccgtg | cccgatgtac | accttccaga | tgtacagctc | 2460 |

| | | | | | | |
|------------|-------------|-------------|------------|------------|------------|------|
| ccgaaagtcc | cagagatgaa | gctcccgaag | gtgcccgaga | tggccgtgcc | cgatgtacac | 2520 |
| cttcagatg | tacagctccc | gaaagttcca | gagatgaagc | taccagagat | gaagctcccg | 2580 |
| aaggtgccgg | agatggccgt | gccggatgta | cgactcccgg | aagttcagct | gcccaaagtg | 2640 |
| tctgaggtga | agctcccaaa | gatgcctgag | atggccgtgc | ctgatgtcca | cctcccggag | 2700 |
| ctacaacttc | ccaaaatgtc | cgaggtgaag | ctcccaaaga | tgcccagat | ggccgtgccc | 2760 |
| gatgttcgcc | tcccgggaagt | tcagctgccc | aaagtgtcag | agatgaaact | ccctaagatg | 2820 |
| ccagagatga | ccatgcccg | cattcgcctc | ccagaagttc | agttgcccaa | agtgcctgac | 2880 |
| attaaacttc | ctgaaatgaa | gcttcacaga | ataaaactcc | ccaaagtgcc | tgacatggca | 2940 |
| gtgcctgatg | tcccccttcc | agagctgcag | ctgcccaaag | tgtcggacat | tccggtgcct | 3000 |
| gaaatgcaag | tgtcacaggt | cccagaggtg | cagcttccca | agatgccaga | gatgaagttg | 3060 |
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| tttagtttca | agttgcccaa | gatgaccatg | ccaagttgg | ggaaagtggg | caagcctggg | 3180 |
| gaggcaagta | ttgaggttcc | agacaaactc | atgacacttc | cctgtctgca | gccagaggtg | 3240 |
| ggcactgagg | catcccatgt | tgggtgtccct | tccctctctc | tccctctgt | ggagcttgac | 3300 |
| ttgcctgggg | ccctgggcct | ggagggacaa | gtccaagaag | ctgtcccagg | caaagtggag | 3360 |
| aagccagagg | gccccaggg | agcagtgggt | gttgagagg | tgggctttcg | tgtgccctct | 3420 |
| gtggagattg | tactctctca | gctgccca | gttgaagttg | agaaagagca | gctagagatg | 3480 |
| gtggagatga | aagtcaaacc | ctcttccaag | ttctctctgc | ccaaattcgg | actttcaggg | 3540 |
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| aagtttacca | tctcacttcc | caaagctcga | gcagggactg | aggccgaagc | gaagggagct | 3660 |
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| atgccatctt | ttgggttg | ccgaggaaag | gaagcagaaa | ctcaggatgg | acgtgtcagc | 3960 |
| cccggggaaa | agctggaggc | catagctggg | cagcttaaga | tccctgcgg | ggaattgg | 4020 |
| acaccgggag | ctcaggagac | agagaagg | accagtggag | tgaagccgtc | aggcctccag | 4080 |
| gtgtccacca | ctgggcaggt | ggttgccagag | ggccaggaga | gtgtgcagag | ggtgtccaca | 4140 |
| ctaggtatct | ctttgcccc | ggtggaattg | gccagctttg | gggaggcagg | ccctgagatc | 4200 |
| gtagccctt | ctgcagagg | cacagcaggc | tctagggctc | aggtgccaca | ggtgatgctg | 4260 |
| gagctacctg | gaaccaggt | ggcaggggg | gatctgttag | tgggtgagg | catcttcaag | 4320 |

| | | | | | | |
|-------------|------------|-------------|-------------|------------|-------------|------|
| atgcccacag | tgacagtgcc | ccagctagag | ctggatgtgg | ggctgggcca | tgaagcccag | 4380 |
| gctggtgaag | cagccaagag | tgaggggtggg | ataaagttga | agttgcccac | actggggacc | 4440 |
| ggaagcagag | gagagggcgt | tgagccccag | ggccccgagg | cccagcgcac | cttcacctc | 4500 |
| tcattgcccg | atgtggaact | cacgtcacca | gtgagtagcc | acgctgagta | ccaggtagtt | 4560 |
| gaggggtgatg | gggatgggtg | gcacaaactc | aagggttcggc | tgcccctggt | tgggtctggca | 4620 |
| aaggccaagg | aagggataga | agttggagaa | aaggttaaga | gtccaaagct | caggctaccc | 4680 |
| cgagtgggct | tcagccagag | tgagtcggtc | tccggagaag | gctctccaag | tcctgaggag | 4740 |
| gaggaagaag | gcagtgggga | aggggcttcc | agtcgccggg | gtagggtaag | ggtccgcctg | 4800 |
| cctcgggtag | gcttggcttc | cccttctaaa | gtctctaagg | gacaggaggg | tgatgcaacc | 4860 |
| tccaagtccc | cagttgggga | gaagtcaccc | aaattccgtt | ttcctagggt | gtccttaagc | 4920 |
| cccaaggccc | ggagtgggag | tagggaccgg | gaagaagggtg | gattcagggt | ccgactgccc | 4980 |
| agtgtgggat | tttcagaaac | agcagttcca | ggttccacca | ggattgaggg | aaccaggct | 5040 |
| gctgccatct | gaagccccag | gacagctgtg | gattccccct | cttgtctttc | cattccccag | 5100 |
| cctagccccc | cattttgtgt | gtgacattac | tagcactaat | cctcagaggg | cttgaagggtg | 5160 |
| agtaactgac | tcaggcagga | gccagtggcc | tgtgccacct | cattggccaa | agtgcctgta | 5220 |
| tatcatgtca | aactatggga | ataaaataat | tcaaaagttg | tcatgtgtct | tggttctcgt | 5280 |
| gggggacaca | aggtctcttt | atgtttcctt | catctggctt | gtgcagtgtt | acctcagctt | 5340 |
| gaacttaaaa | tcttgagcc | ttgggggctg | gagaggtggc | ccagaggtta | agagcactgg | 5400 |
| ctgctcatgc | agaggtcctg | agttcaattc | cc | | | 5432 |

<210> 30
 <211> 468
 <212> DNA
 <213> Arabidopsis

| | |
|-------------|------------|
| <400> 30 | |
| tttaagattg | tgggatgggc |
| ttcaaagtag | ctgggccttc |
| tcttcgctta | aggagttccg |
| ggaacttttg | ggaattccgg |
| caacttgga | agttgaggca |
| tcgtcggtag | ctctggcttt |
| ggaatctctg | gaactgcagg |
| ttttgtaatt | tctggaacct |
| tgggcagttc | aggttttgga |
| acttctggaa | tcttcggcag |
| ttcaggcttt | ggaatctctg |
| gaaacttagg | agcctcagac |
| tttggaacct | ctggaaccgt |
| tggcaattcg | ggctttggaa |
| cctctggcac | cttcggcaac |
| tcaggcttct | gaatctccgg |
| catcttgggc | aactcgggtt |
| tcttgaatct | ccggcatctt |
| aggtaactcc | ggcatttttg |
| gcaactcccg | gcttctggat |
| ctccgggtacc | ttaagagcct |
| caagcttggg | aactcttggc |
| aactttggca | actcaaggct |
| tttggaat | |

<210> 31
 <211> 668
 <212> DNA
 <213> Mouse

<220>
 <221> misc_feature
 <222> (1)..(668)
 <223> n equals unknown

<400> 31
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 attagtgcta gtaatgtcac acacaaaatg gggggctagg ctggggaatg gaaagacaag 180
 agggggaatc cacagctgtc ctggggcttc agatggcagc agcctggggtt ccctcaattc 240
 ctggtggaac ctggaactgc tgtttctgaa aatcccacac tgggcagtcg gaccctgaat 300
 ccaccttctt cccggtccct actcccactc cgggccttgg ggcttaagga caccctagga 360
 aaacggaatt tgggtgactt ctccccaaact ggggacttgg aggttgcac accctcctgt 420
 ccctagagac tttagaaggg gaagccaagc ctacccgagg caggcggacc cttaccctac 480
 cccggcgact ggaagccctt tcccactgc cttcttcctc ctccctaggac ttggagagcc 540
 ttctcccgag accgactcac tctggcttga acccactcng ggtagcctga gctttgactc 600
 ttagcctttt ctcccacttc tatcccttcc ttgccctttg cagaccaaca agggaggcgg 660
 accttgag 668

<210> 32
 <211> 433
 <212> DNA
 <213> Mouse

<220>
 <221> misc_feature
 <222> (1)..(433)
 <223> n equals unknown

<400> 32
 cagaaactca gatggacgtg tcagccccgg ggaaaagctg gaggccatag ctgggcagct 60
 taagatccct gcggtggaat tggtcacacc gggagctcag gagacagaga aggtcaccag 120
 tggagtgaag ccgtcaggcc tcaggtgtc caccactggg caggtggttg cagagggcca 180
 ggagagtgtg cagaggggtgt ccacactagg tatctctttg cccaggtgg aattggccag 240
 ctttggggag gcaggccctg agatcgtagc cccttctgca gagggcacag caggctctag 300

ggtccaggtg ccacaggtga tgctggagct acctggaacc cacgtggcag ggggtgatct 360
 gttagtgggt gagggcatct tcangatgcc ccccgtagaca gtgcccctcc ttgtgctggc 420
 tgtggggctg gcc 433

<210> 33
 <211> 451
 <212> DNA
 <213> Human

<400> 33
 acttccagat tattttattc acatggcttg gtggggtaca ggcactcctg ccagagagac 60
 aggagcaggc ctccctgcc a gccctgggtca gtcacccacc tcccggccct cttaggggtta 120
 gtgctagtta tcacacacac aacagcgagg gggtagagaa aggaaggcaa gaagggatcc 180
 ccatctgact aggggcttca tacagccgca gcctgagccc cctccatcct ggccgggcct 240
 ggagcccctg tctctgaaaa cccacgctg ggcagccgca cccgcaatcc accctcttcc 300
 tgggtcccac tccactccg ggccttgggg cttagggaca ccctggggaa gcggaacttg 360
 ggtgacttct ctctgactgg ggacttgggg gctgcatcgc cctcctgccc ccgagaggct 420
 ttagaagggg ccgccaggcc tacacgtggc a 451

<210> 34
 <211> 711
 <212> DNA
 <213> Mouse

<220>
 <221> misc_feature
 <222> (1)..(711)
 <223> n equals unknown

<400> 34
 gcacgtgag tacaggtagt gaggggtgatg ggatgggtgg cacaaactca aggttcggct 60
 gccctgttg gtctgcanaa ggccaagnga gggatagaag tggagaaaag gctaagagtc 120
 caaagctcag gctaccccga gtgggcttca gccagagtga gtcggtctcc ggagaaggct 180
 ctccaagtcc tgaggaggag gaagaaggca gtggggaagg ggcttccagt cgccggggta 240
 gggtaagggt ccgctgcct cgggtaggct tggcttcccc ttctaaagtc tctaaggagc 300
 aggaggggtga tgcaacctcc aagtccccag ttggggagaa gtcacccaaa ttccgttttc 360
 ctaggggtgc cttaagcccc aaggcccga gtgggagtag ggaccgggaa gaagggtgat 420
 tcagggtccg actgcccagt gtgggatttt cagaaacagc agttccaggt tccaccagga 480
 ttgagggaac ccaggctgct gccatctgaa gccccaggac agctgtggat tccccctctt 540
 gtctttccat tccccagcct agccccccat tttgtgtgtg acattactag cactaatcct 600

cagagggctt gaaggtgggt aactgactca ggcaggagcc agtggcctgt gccacctcat 660
 tggccaaagt gcctgtatat catgtcaaac tatgggaata aaataattca a 711

<210> 35
 <211> 239
 <212> DNA
 <213> Mouse

<220>
 <221> misc_feature
 <222> (1)..(239)
 <223> n equals unknown

<400> 35
 ggattggggg naccaggct cctgccttct gaacccccag gacacctgtg gattccccct 60
 catgtctttc cattccccac cctaccccc catttttgtgt gtgacattac taccactagt 120
 cctcagaggg cttgaagggt ggtaactgac tcaggcagga ccagtgccc tgtgccacct 180
 cattggccta agtgcctgta tttcatgtca aactatggga ataaaataat tctaaagtt 239

<210> 36
 <211> 290
 <212> DNA
 <213> Mouse

<400> 36
 aacttcccca tgccccactt tccgacatcc agacaccgca gttccacgtt ccaccagcac 60
 agagggcacc cagggcgccc caatctggag cccaggaca cctctggatt cccctctag 120
 tctttccatt cccacccta cccccccatt ttgtgtgtga cattactagc actattcctc 180
 agagggcttg aaggtgggta actgactcag gcaggagcca gtggcctgtc ccacctcatt 240
 gcccaaagtg cctgtatatc atgtcaaact atgggaataa aataattcaa 290

<210> 37
 <211> 326
 <212> DNA
 <213> Mouse

<400> 37
 accatacaaa gaataccctc ctggacgaac aagccttcac ggtccccccc cccaatctgc 60
 gattttcagc cacaccagtt ccagttccac caggattgag ggaaccacc ctgctgcctt 120
 atggatcccc aggacagttg tggattcccc cttttgtctt tccattcccc agccttcccc 180
 cccattttgt gtgtgacatt actaccacta atcctcagag ggcttgaagg tgggtaactg 240
 actcaggcag gaccagtggt cctgtgccac ctcatgggcc aaagtgcctg tatttcatgt 300
 caaactatgg gaataaaata attcaa 326

<210> 38
 <211> 633
 <212> DNA
 <213> Mouse

<400> 38
 acgccatggc gagctctggg cgagttgccg tcttgagact ccatggcgac cccttgctgc 60
 ccctcgcagg agctgagacg ggcggagttg gtggagatta tcgtggagac cgaggcacag 120
 accgggggtca gcggcttcaa cgtagcaggc ggcggcaaag aaggaatctt tgtccgtgag 180
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 agtgcccgtg tggtctttga gaacttcaaa tatgaggatg cacttcgcct gctgcaatgc 300
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 ctgaggcccc ggacggtgtc tggatacgag atgaagggcc cacgggcaa agtggccaag 420
 ctgaacatcc agagtctggc ccctgtgaag aagaagaaga tggtagactg ggccctgagg 480
 acccctgcag atttggcccc tgttgacgtc gagttctctt tttccaagtt ctcccgactg 540
 gcgtgggggtc ttcaaagccg agctgtcaaa ggacctggtc cagctggccc aaccgctcgc 600
 ccgcttcagc tgccctcgggt ttgggtccca gaa 633

<210> 39
 <211> 488
 <212> DNA
 <213> Human

<400> 39
 tttacttcca gattatttta ttcacatggc ttggtggggt acaggcactc ctgccagaga 60
 gacaggagca ggcctccctg ccagccctgg tcagtcaccc acctcccggc cctcttaggg 120
 ttagtgctag ttatcacaca cacaacagcg agggggtaga gaaaggaagg caagaaggga 180
 tccccatctg actagggggt tcagacagcc gcagcctgag cccctccat cctggccggg 240
 cctggagccc ctgtctctga aaaccccacg ctgggcagcc gcacccgcaa tccaccctct 300
 tcctggtccc cactcccact ccgggccttg gggcttaggg acaccctggg gaagcggaac 360
 ttgggtgact tctctctgac gggggacatt ggggctgcat cgccctcctg ccccgagag 420
 gctttagaag gggccgccag gcctacacgt ggcaagcgga cccggacccg gccccggga 480
 cccgaggc 488

<210> 40
 <211> 372
 <212> DNA
 <213> Human

```

<400> 40
tttactttcca gattatttta ttcacatggc ttggtggggt acaggcactc ctgccagaga      60
gacaggagca ggctccctg ccagccctgg tcagtcaccc acctcccggc cctcttaggg      120
ttagtgctag ttatcacaca cacaacagcg agggggtaga gaaaggaagg caagaaggga      180
tccccatctg actaggggct tcagacagcc gcagcctgag ccccttcat tctgggcccgg      240
ccttgaaccc ctggtttttg aaaacccaac cttggccagc cgaccccgaa atcaaccctt      300
ttcttgggtcc caactccac tccgggcctt ggggcttagg gacaccctgg ggaagcggaa      360
cttgggtgac tt                                                                372

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<210> 41
<211> 642
<212> DNA
<213> Mouse

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<220>
<221> misc_feature
<222> (1)..(642)
<223> n equals unknown

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```

<400> 41
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aggggtggcac aggggccact ggcttcctgc ctgagtcagt taccgccnnac cttcaagccc      120
tctgagggat tagtgctagt naatgtcaca cacaanaatg gggggctagg nctggggaat      180
ggaaagacaa gagggggaat ccacagctgt cctggggcctt cagatggcag cagcctgggt      240
tccctcaatc ctggtggaac ctggaactgc tgtttctgaa aatcccacac tgggcagtcg      300
gaccctgaat ccaccttctt ccccgctccc tacctccact cccggccttg ggggcttaag      360
gacacccta gaaaacggga atttgggtga cttctcccca actggggact tggagggttg      420
atcacctcc tgtcccttag agactttaga aggggaagcc aagcctaccc gaagcaggcg      480
gacccttacc ctacccccgc gactggaagc cccttagcca ctgctcttcc tcctcctagg      540
actggaaagc cctcttcgag accgactacc ttggctgaaa ccactcggg gtagctgagc      600
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<210> 42
<211> 328
<212> DNA
<213> Mouse

```

```

<400> 42
cccacgcgtc cgggagatga aagtcaaacc ctcttccaag ttctctctgc ccaaattcgg      60
actttcaggg cccaaagctg tcaagggaga ggtggagggg cctgggcgag ccaccaagct      120

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gaagggtttcc aagtttacca tctcacttcc caaagctcga gcagggactg aggccgaagc 180
gaagggagct ggggaagccg ggttgctgcc agccctggat ctgtccatcc cacagctcag 240
cctggatgcc cagctgcctt caggcaagggt ggaagtagct gatagcaagc ctaaatcgtc 300
cagatttgct ctgcccaagt ttgggggt 328

<210> 43
<211> 564
<212> DNA
<213> Mycosphaerella graminicola

<400> 43
cgctcccggc ttgacaact cccgataccc cgacactctc gaccctaag cttccatctc 60
tcagcaccac tagcattaag ctgcgcgcgt tctctgctcc tacactcccg gcttttacga 120
tccctaact cccgacactc tcgaccccta agcttccatc tctcagcacc actagcatta 180
agctgcgcgc gttctctgct cctacactcc cggcctttac gatccctaac ctcccgatat 240
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cgggcctccc caggttcacc acgagcagca tcaacttgcc gtcgttctcg gctccaagcc 480
tcccgggcct cccacgttc accacgaaca gcgtcaactt gccgtcgttc tcgggtccaa 540
gcctcccggc cctccccacg ttca 564

<210> 44
<211> 238
<212> DNA
<213> Mouse

<220>
<221> misc_feature
<222> (1)..(238)
<223> n equals unknown

<400> 44
gaatgaggga acccagactt ntgccttttg aagcccccg acagtcgtgg gttccccttc 60
tagtctttcc tttcccagc ctagcccccc attttgctgtg tgacattact agcactattc 120
ctcagagggt ttgaagggtg gtaattgact caggcaggag ccagtggcct gtgccacctc 180
attggccaaa gtgcctgtat atcatgtcaa actatgggaa tgaaattatt caaaagtt 238

<210> 45
<211> 654
<212> DNA
<213> Mouse

<400> 45
aaggttaggc tgcccatggt tggctctggca aaggccaagg aagggataga agttggagaa 60
aaggctaaga gtccaaagct caggctaccc cgagtgaact tcagccagag tgagtcggtc 120
tccggagaag gctctccaag tcctgaggag gaggaagaag gcagtgggga aggggcttcc 180
agtcgccggg gtagggtaag ggtccgcctg cctcgggtag gcttggttc cccttctaaa 240
gtctctaagg gacaggaggg tgatgcaacc tccaagtccc cagttgggga gaagtcaccc 300
aaattccgtt ttcctagggg gtccttaagc cccaaggccc ggagtgggag tagggaccgg 360
gaagaaggtg gattcagggt ccgactgccc agtgtgggat tttcagaaac agcagttcca 420
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<211> 232
<212> DNA

<213> Human

<400> 47

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ccttgggggt gatggacacc ctgggggaaac ggaacttggg tgaattcttt ttgacagggg      180
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<211> 418

<212> DNA

<213> Mouse

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ggattagtgc tagtaatgtc acacacaaaa tgggggggcta ggctggggaa tggaaagaca      180
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<210> 49

<211> 451

<212> DNA

<213> Human

<400> 49

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gtgctagtta tcacacacac aacagcgagg gggtagagaa aggaaggcaa gaagggatcc      180
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<211> 498

<212> DNA

<213> Mouse

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<210> 51
<211> 409
<212> DNA
<213> Mouse

<400> 51
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<211> 356
<212> DNA
<213> Mouse

<400> 52
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<210> 53

<211> 356
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 <213> Mouse

<400> 53
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 <211> 73
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 <213> Mouse

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<210> 55
 <211> 342
 <212> DNA
 <213> Mouse

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<210> 56
 <211> 390
 <212> DNA
 <213> Human

<400> 56
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 gcaagccagg cgaggcgggt gctgagggtc cagggaaagt ggtaacactt ccctgtctgc 180
 agccagaggt ggatgggtgag gctcatgtgg gtgtccctct ctactctgc cttcagtgga 240

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<210> 57
 <211> 5432
 <212> DNA
 <213> Mouse

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| cttacagctg tgcatacccc cgagtggggc tctgtcagg agaaaaggcc atcactcaag | 180 |
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| gcagcgctga ggagctgaga cgggcggagt tggtgagat tatcgtggag accgaggcac | 360 |
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| tgagtgcccg tgtgttcttt gagaacttca aatatgagga tgacttcgc ctgctgcaat | 540 |
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| cttccatgcc | tggacaccca | ggaaggagct | gcagtggtaa | aagtccctac | cctggatgtg | 1860 |
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| gaggttcctg | aagtggccct | caagatgccc | cggctcagtt | tccccgttt | tgggattcgg | 1980 |
| gggaaggaag | ccactgaagc | caaagtagtc | aagggcagcc | ctgaggccaa | agcaaaggg | 2040 |
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 <213> Mouse

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 <213> Mouse

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 <211> 494
 <212> DNA
 <213> Mouse

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 <223> n equals unknown

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 <212> DNA
 <213> Mouse

<220>
 <221> misc_feature
 <222> (1)..(360)
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<210> 62
 <211> 359
 <212> DNA
 <213> Mouse

<400> 62
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<210> 63
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 <212> DNA
 <213> Rat

<400> 63
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| tccagggtccc gcaagtggaa ctccccaccc tgccctcttt accactctg cccacacttc | 1200 |
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aaagttaaaa aaaaaaaaaa a 4641

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```

<210> 64
<211> 672
<212> DNA
<213> Human

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<220>
<221> misc_feature
<222> (1)..(672)
<223> n equals unknown

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```

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<210> 65
<211> 597
<212> DNA
<213> Mouse

```

```

<400> 65
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<210> 66
<211> 697
<212> DNA
<213> Human

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<220>
<221> misc_feature
<222> (1)..(697)
<223> n equals unknown

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<400> 66
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<210> 67
<211> 626

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<212> DNA
<213> Human

<220>
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<222> (1)..(626)
<223> n equals unknown

<400> 67
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<210> 68
<211> 645
<212> DNA
<213> Human

<400> 68
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421

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| agctgcccaa aatgtcagac atgaaacttc caaagatccc tgagatggct gtaccgatg | 1740 |

| | | | | | | |
|-------------|------------|-------------|-------------|-------------|------------|------|
| ttcaccttcc | ggaagtgaag | ctgccccaaag | tccccgagat | gaaagtccca | gaaatgaagc | 1800 |
| ttccgaagat | cccggagatg | gccgtgcctg | atgtacacct | tccagatata | cagctcccga | 1860 |
| aagttcccga | gatgaagctc | ccagacatga | agctccccga | ggtgcctgag | atggccgtgc | 1920 |
| ctgatgtaca | ccttccagat | atacagctcc | cgaaagttcc | cgagatgaag | ctcccagaca | 1980 |
| tgaagctccc | gaaggtgcct | gagatggccg | tgcttgatgt | acgaattccg | gaagttcagc | 2040 |
| tacccaaagt | gtccgaggtg | aagctcccga | agataccgga | catggccgtg | cctgatgttc | 2100 |
| gcctcccaga | gctgcaactg | cccaaaatgt | ctgaggtgaa | gctcccgaag | ataccggaca | 2160 |
| tgggccgtacc | tgatgttcgc | ctcccagaag | ttcagctacc | caaagtgtca | gagctgaagc | 2220 |
| tcccgaaggt | gcctgagatg | accatgcccg | acattcgcct | cccggaagtt | cagctgccc | 2280 |
| aagtgctga | cattaaactt | ccagaaataa | aactccccaa | agtgctgag | atggccgtgc | 2340 |
| ctgatgtccc | ccttccagaa | ctacagctgc | ccaaagtgcc | acaggtccca | gacgtgcatc | 2400 |
| ttcccaaagt | gccagagatg | aagttgccc | aggttcctga | ggcacagagg | aaatctgcag | 2460 |
| gggcgagca | ggcagaaaag | accgaattta | gcttcaagtt | gccaagatg | actgtgccc | 2520 |
| agttggggaa | agtgaccaag | cctggggagg | caggtattga | ggttccagac | aaactcctga | 2580 |
| tacttcctg | tctgcagcca | gaggtgggca | ctgaggtggc | ccgtgttgg | gtcccttccc | 2640 |
| tctctctccc | ttctgtggag | cttgacttgc | ctggggccct | gggcctggag | ggacaagtcc | 2700 |
| aagaagctgt | ctctggcaaa | gtggagaagc | cagagggccc | caggggtggca | gtagggactg | 2760 |
| gagaggcggg | cttcgcgtg | ccctctgtgg | agattgtcaa | tcctcagctg | cccacggttg | 2820 |
| aagtcaagaa | agagcagcta | gagatggtgg | agatgaaagt | caaaccact | tccaagttct | 2880 |
| ctctgccc | atttgactt | tcagggccc | aagctgtcaa | ggcagaggtg | gaggggctg | 2940 |
| ggcgagccac | caagctgaag | gtatccaagt | ttgccatctc | gcttcccaga | gctcgagcag | 3000 |
| ggactgacgc | ggacgcgaag | ggagctgggg | aagcggggtt | gctgcctgcc | ctcgatctgt | 3060 |
| ccatcccaca | gctcagcctg | gatgetcaac | tgccctcagg | caaggtggag | gtagcagggg | 3120 |
| ctgagagcaa | gcctaaaggg | tccagatttg | ctctgccc | gtttggggcg | aaaggccggg | 3180 |
| actctgaagc | cgacgtactg | gtggcagggg | aggctgagct | ggaggggaag | ggttggggct | 3240 |
| gggacgggaa | ggtgaagatg | ccaagctga | agatgccatc | ttttgggctg | tcccaggaa | 3300 |
| aagaagcaga | aattcaggat | gggcgtgtca | gcccaggaga | aaagctggaa | gccatagctg | 3360 |
| ggcagcttaa | gatccctgag | gtggaactgg | tcacaccagg | agctcaggag | acagagaagg | 3420 |
| tcaccagtgg | agtgaagcca | tcaggcctcc | agggtgtccac | cactaggcag | gtggttgag | 3480 |
| agggccagga | gggggcgcag | aggggtgtcct | cattaggtat | ctctttgccc | caggtggaac | 3540 |

tggccagctt tggggaggca ggccttgaga tcgcagcccc atctgcagag ggcacagtag 3600
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 gtgatctgtt agtgggtgag ggcattctca agatgccac agtgacagtg cccagtttag 3720
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 agagccccga ggcccagcac accttcaca tctcattgcc tgacgtagaa ctcacatcac 3900
 cagtgagtag ccacgctgag taccaggtgg ttgagggcga tggggatggc gggcacaaac 3960
 tcaaggtgag gctgcccctg tttggtctgg caagggccaa ggaaggaata gaaactggag 4020
 aaaaggttaa aagtccaaag ctcaggctac cccgagtggg cttcagccaa agtgagtcgg 4080
 cctctggaga aggctctccc agtcctgagg aggaggaaga aggcagtggg gaaggggctt 4140
 ccggtcgccg tggtcgggtc agggctcgct tgcctcgtgt aggccttggt tccccttcta 4200
 aaggctctaa gggacaggag ggtgatgcgg cctccaagtc cccagttggg gagaagtccc 4260
 ccaagttccg ctttcttagg gtgtccttaa gcccgaaggc ccggagtggg agtaaggacc 4320
 gggaagaagg tggattcagg gtccgactgc ccagtgtggg attttcagaa acagcagctc 4380
 caggctccgc caggattgag gggaccagc ctgctgccat ctgaagccct gggacagctg 4440
 tggattcccc ctcttgtctt cccatcccca tccctgctcc ccattttatg tgtgacatta 4500
 ctgactacta tctcagagg gcttgaaggt gggcagctga ctcaggcagg agcgggtctgt 4560
 gccacctcat tggctgacgt gcctgtatat catgccaagc tctgtgaata aaataattca 4620
 aaagttaaaa aaaaaaaaaa a 4641

<210> 78
 <211> 370
 <212> PRT
 <213> Arabidopsisthaliana

<400> 78

Met Ala Leu Met Lys Lys Ser Leu Ser Ala Ala Leu Leu Ser Ser Pro
 1 5 10 15

Leu Leu Ile Ile Cys Leu Ile Ala Leu Leu Ala Asp Pro Phe Ser Val
 20 25 30

Gly Ala Arg Arg Leu Leu Glu Asp Pro Lys Pro Glu Ile Pro Lys Leu
 35 40 45

Pro Glu Leu Pro Lys Phe Glu Val Pro Lys Leu Pro Glu Phe Pro Lys
 50 55 60

Pro Glu Leu Pro Lys Leu Pro Glu Phe Pro Lys Pro Glu Leu Pro Lys
65 70 75 80

Ile Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Val Pro Glu Ile Pro
85 90 95

Lys Pro Glu Glu Thr Lys Leu Pro Asp Ile Pro Lys Leu Glu Leu Pro
100 105 110

Lys Phe Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Met Pro Glu Ile
115 120 125

Pro Lys Pro Glu Leu Pro Lys Val Pro Glu Ile Gln Lys Pro Glu Leu
130 135 140

Pro Lys Met Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Phe Pro Glu
145 150 155 160

Ile Pro Lys Pro Asp Leu Pro Lys Phe Pro Glu Asn Ser Lys Pro Glu
165 170 175

Val Pro Lys Leu Met Glu Thr Glu Lys Pro Glu Ala Pro Lys Val Pro
180 185 190

Glu Ile Pro Lys Pro Glu Leu Pro Lys Leu Pro Glu Val Pro Lys Leu
195 200 205

Glu Ala Pro Lys Val Pro Glu Ile Gln Lys Pro Glu Leu Pro Lys Met
210 215 220

Pro Glu Leu Pro Lys Met Pro Glu Ile Gln Lys Pro Glu Leu Pro Lys
225 230 235 240

Leu Pro Glu Val Pro Lys Leu Glu Ala Pro Lys Val Pro Glu Ile Gln
245 250 255

Lys Pro Glu Leu Pro Lys Met Pro Glu Leu Pro Lys Met Pro Glu Ile
260 265 270

Gln Lys Pro Glu Leu Pro Lys Met Pro Glu Ile Gln Lys Pro Glu Leu
275 280 285

Pro Lys Val Pro Glu Val Pro Lys Pro Glu Leu Pro Thr Val Pro Glu
290 295 300

Val Pro Lys Ser Glu Ala Pro Lys Phe Pro Glu Ile Pro Lys Pro Glu
305 310 315 320

Leu Pro Lys Ile Pro Glu Val Pro Lys Pro Glu Leu Pro Lys Val Pro
325 330 335

Glu Ile Thr Lys Pro Ala Val Pro Glu Ile Pro Lys Pro Glu Leu Pro
340 345 350

Thr Met Pro Gln Leu Pro Lys Leu Pro Glu Phe Pro Lys Val Pro Gly
355 360 365

Thr Pro
370

<210> 79
<211> 1389
<212> PRT
<213> Rat

<400> 79

Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
1 5 10 15

Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Phe
20 25 30

Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg
35 40 45

Glu Asp Ser Pro Ala Ala Lys Ser Leu Ser Leu Gln Glu Gly Asp Gln
50 55 60

Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala
65 70 75 80

Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu
85 90 95

Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val
100 105 110

Ser Gly Tyr Glu Met Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Asn
115 120 125

Ile Gln Ser Leu Ser Pro Val Lys Lys Lys Lys Met Val Ile Gly Thr
130 135 140

Leu Gly Thr Pro Ala Asp Leu Ala Pro Val Asp Val Glu Phe Ser Phe
145 150 155 160

Pro Lys Phe Ser Arg Leu Arg Arg Gly Leu Lys Ala Asp Ala Val Lys
165 170 175

Gly Pro Val Pro Ala Ala Pro Ala Arg Arg Arg Leu Gln Leu Pro Arg
180 185 190

Leu Arg Val Arg Glu Val Ala Glu Glu Ala Gln Val Ala Arg Met Ala
195 200 205

Ala Ala Ala Pro Pro Ser Arg Lys Ala Lys Ser Glu Ala Glu Val Ala
210 215 220

Thr Gly Ala Gly Phe Thr Ala Pro Gln Ile Glu Leu Val Gly Pro Arg
225 230 235 240

Leu Pro Ser Ala Glu Val Gly Val Pro Lys Val Ser Val Pro Lys Gly
245 250 255

Thr Pro Ser Thr Glu Ala Ala Ser Gly Phe Ala Leu His Leu Pro Thr
260 265 270

Leu Gly Leu Gly Ala Pro Ala Ala Pro Ala Val Glu Pro Pro Thr Thr
275 280 285

Gly Ile Gln Val Pro Gln Val Glu Leu Pro Thr Leu Pro Ser Leu Pro
290 295 300

Thr Leu Pro Thr Leu Pro Cys Leu Asp Thr Gln Glu Gly Ala Ala Val
305 310 315 320

Val Lys Val Pro Thr Leu Asp Val Ala Ala Pro Ser Val Glu Val Asp
325 330 335

Leu Ala Leu Pro Gly Ala Glu Val Glu Ala Gln Gly Glu Val Pro Glu
340 345 350

Val Ala Leu Lys Met Pro Arg Leu Ser Phe Pro Arg Phe Gly Val Arg
355 360 365

Gly Lys Glu Ala Thr Glu Ala Lys Val Val Lys Gly Ser Pro Glu Ala
370 375 380

Lys Ala Lys Gly Pro Arg Leu Arg Met Pro Thr Phe Gly Leu Ser Leu
 385 390 395 400

Leu Glu Ser Arg Pro Ser Gly Pro Glu Val Ala Ala Glu Ser Lys Leu
 405 410 415

Lys Leu Pro Thr Leu Lys Met Pro Ser Phe Gly Ile Ser Val Ala Gly
 420 425 430

Pro Glu Val Lys Ala Pro Lys Gly Pro Glu Val Lys Leu Pro Lys Val
 435 440 445

Pro Glu Ile Lys Leu Pro Lys Ala Pro Glu Ala Ala Ile Pro Asp Val
 450 455 460

Gln Leu Pro Glu Val Gln Leu Pro Lys Met Ser Asp Met Lys Leu Pro
 465 470 475 480

Lys Ile Pro Glu Met Ala Val Pro Asp Val His Leu Pro Glu Val Lys
 485 490 495

Leu Pro Lys Val Pro Glu Met Lys Val Pro Glu Met Lys Leu Pro Lys
 500 505 510

Ile Pro Glu Met Ala Val Pro Asp Val His Leu Pro Asp Ile Gln Leu
 515 520 525

Pro Lys Val Pro Glu Met Lys Leu Pro Asp Met Lys Leu Pro Lys Val
 530 535 540

Pro Glu Met Ala Val Pro Asp Val His Leu Pro Asp Ile Gln Leu Pro
 545 550 555 560

Lys Val Pro Glu Met Lys Leu Pro Asp Met Lys Leu Pro Lys Val Pro
 565 570 575

Glu Met Ala Val Pro Asp Val Arg Ile Pro Glu Val Gln Leu Pro Lys
 580 585 590

Val Ser Glu Val Lys Leu Pro Lys Ile Pro Asp Met Ala Val Pro Asp
 595 600 605

Val Arg Leu Pro Glu Leu Gln Leu Pro Lys Met Ser Glu Val Lys Leu
 610 615 620

Pro Lys Ile Pro Asp Met Ala Val Pro Asp Val Arg Leu Pro Glu Val
 625 630 635 640
 Gln Leu Pro Lys Val Ser Glu Leu Lys Leu Pro Lys Val Pro Glu Met
 645 650 655
 Thr Met Pro Asp Ile Arg Leu Pro Glu Val Gln Leu Pro Lys Val Pro
 660 665 670
 Asp Ile Lys Leu Pro Glu Ile Lys Leu Pro Lys Val Pro Glu Met Ala
 675 680 685
 Val Pro Asp Val Pro Leu Pro Glu Leu Gln Leu Pro Lys Val Pro Gln
 690 695 700
 Val Pro Asp Val His Leu Pro Lys Val Pro Glu Met Lys Leu Pro Lys
 705 710 715 720
 Val Pro Glu Ala Gln Arg Lys Ser Ala Gly Ala Glu Gln Ala Glu Lys
 725 730 735
 Thr Glu Phe Ser Phe Lys Leu Pro Lys Met Thr Val Pro Lys Leu Gly
 740 745 750
 Lys Val Thr Lys Pro Gly Glu Ala Gly Ile Glu Val Pro Asp Lys Leu
 755 760 765
 Leu Ile Leu Pro Cys Leu Gln Pro Glu Val Gly Thr Glu Val Ala Arg
 770 775 780
 Val Gly Val Pro Ser Leu Ser Leu Pro Ser Val Glu Leu Asp Leu Pro
 785 790 795 800
 Gly Ala Leu Gly Leu Glu Gly Gln Val Gln Glu Ala Val Ser Gly Lys
 805 810 815
 Val Glu Lys Pro Glu Gly Pro Arg Val Ala Val Gly Thr Gly Glu Ala
 820 825 830
 Gly Phe Arg Val Pro Ser Val Glu Ile Val Asn Pro Gln Leu Pro Thr
 835 840 845
 Val Glu Val Lys Lys Glu Gln Leu Glu Met Val Glu Met Lys Val Lys
 850 855 860
 Pro Thr Ser Lys Phe Ser Leu Pro Lys Phe Gly Leu Ser Gly Pro Lys

| | | | | | | |
|---|-----|-----|--|------|--|------|
| 865 | | 870 | | 875 | | 880 |
| Ala Val Lys Ala Glu Val Glu Gly Pro Gly Arg Ala Thr Lys Leu Lys | | | | | | |
| | 885 | | | 890 | | 895 |
| Val Ser Lys Phe Ala Ile Ser Leu Pro Arg Ala Arg Ala Gly Thr Asp | | | | | | |
| | 900 | | | 905 | | 910 |
| Ala Asp Ala Lys Gly Ala Gly Glu Ala Gly Leu Leu Pro Ala Leu Asp | | | | | | |
| | 915 | | | 920 | | 925 |
| Leu Ser Ile Pro Gln Leu Ser Leu Asp Ala Gln Leu Pro Ser Gly Lys | | | | | | |
| | 930 | | | 935 | | 940 |
| Val Glu Val Ala Gly Ala Glu Ser Lys Pro Lys Gly Ser Arg Phe Ala | | | | | | |
| 945 | | 950 | | 955 | | 960 |
| Leu Pro Lys Phe Gly Ala Lys Gly Arg Asp Ser Glu Ala Asp Val Leu | | | | | | |
| | 965 | | | 970 | | 975 |
| Val Ala Gly Glu Ala Glu Leu Glu Gly Lys Gly Trp Gly Trp Asp Gly | | | | | | |
| | 980 | | | 985 | | 990 |
| Lys Val Lys Met Pro Lys Leu Lys Met Pro Ser Phe Gly Leu Ser Arg | | | | | | |
| | 995 | | | 1000 | | 1005 |
| Gly Lys Glu Ala Glu Ile Gln Asp Gly Arg Val Ser Pro Gly Glu | | | | | | |
| 1010 | | | | 1015 | | 1020 |
| Lys Leu Glu Ala Ile Ala Gly Gln Leu Lys Ile Pro Glu Val Glu | | | | | | |
| 1025 | | | | 1030 | | 1035 |
| Leu Val Thr Pro Gly Ala Gln Glu Thr Glu Lys Val Thr Ser Gly | | | | | | |
| 1040 | | | | 1045 | | 1050 |
| Val Lys Pro Ser Gly Leu Gln Val Ser Thr Thr Arg Gln Val Val | | | | | | |
| 1055 | | | | 1060 | | 1065 |
| Ala Glu Gly Gln Glu Gly Ala Gln Arg Val Ser Ser Leu Gly Ile | | | | | | |
| 1070 | | | | 1075 | | 1080 |
| Ser Leu Pro Gln Val Glu Leu Ala Ser Phe Gly Glu Ala Gly Pro | | | | | | |
| 1085 | | | | 1090 | | 1095 |
| Glu Ile Ala Ala Pro Ser Ala Glu Gly Thr Val Gly Ser Arg Ile | | | | | | |
| 1100 | | | | 1105 | | 1110 |

| | | | | |
|---------|-----------------|------|---------------------|-----------------|
| Gln Val | Pro Gln Val Met | Leu | Glu Leu Pro Gly Thr | Gln Val Ala |
| 1115 | | 1120 | | 1125 |
| Gly Gly | Asp Leu Leu Val | Gly | Glu Gly Ile Phe Lys | Met Pro Thr |
| 1130 | | 1135 | | 1140 |
| Val Thr | Val Pro Gln Leu | Glu | Leu Asp Val Gly | Leu Gly His Glu |
| 1145 | | 1150 | | 1155 |
| Ala Gln | Ala Gly Glu Thr | Ala | Lys Ser Glu Gly | Gly Leu Lys Leu |
| 1160 | | 1165 | | 1170 |
| Lys Leu | Pro Thr Leu Gly | Ala | Gly Gly Lys Gly | Glu Gly Ala Glu |
| 1175 | | 1180 | | 1185 |
| Ala Gln | Ser Pro Glu Ala | Gln | His Thr Phe His | Ile Ser Leu Pro |
| 1190 | | 1195 | | 1200 |
| Asp Val | Glu Leu Thr Ser | Pro | Val Ser Ser His | Ala Glu Tyr Gln |
| 1205 | | 1210 | | 1215 |
| Val Val | Glu Gly Asp Gly | Asp | Gly Gly His Lys | Leu Lys Val Arg |
| 1220 | | 1225 | | 1230 |
| Leu Pro | Leu Phe Gly Leu | Ala | Arg Ala Lys Glu | Gly Ile Glu Thr |
| 1235 | | 1240 | | 1245 |
| Gly Glu | Lys Val Lys Ser | Pro | Lys Leu Arg Leu | Pro Arg Val Gly |
| 1250 | | 1255 | | 1260 |
| Phe Ser | Gln Ser Glu Ser | Ala | Ser Gly Glu Gly | Ser Pro Ser Pro |
| 1265 | | 1270 | | 1275 |
| Glu Glu | Glu Glu Glu Gly | Ser | Gly Glu Gly Ala | Ser Gly Arg Arg |
| 1280 | | 1285 | | 1290 |
| Gly Arg | Val Arg Val Arg | Leu | Pro Arg Val Gly | Leu Ala Ser Pro |
| 1295 | | 1300 | | 1305 |
| Ser Lys | Gly Ser Lys Gly | Gln | Glu Gly Asp Ala | Ala Ser Lys Ser |
| 1310 | | 1315 | | 1320 |
| Pro Val | Gly Glu Lys Ser | Pro | Lys Phe Arg Phe | Pro Arg Val Ser |
| 1325 | | 1330 | | 1335 |

Leu Ser Pro Lys Ala Arg Ser Gly Ser Lys Asp Arg Glu Glu Gly
1340 1345 1350

Gly Phe Arg Val Arg Leu Pro Ser Val Gly Pro Thr Ala Gln Cys
1355 1360 1365

Gly Ile Phe Arg Asn Ser Ser Ser Ser Ser Ala Arg Ile Glu Gly
1370 1375 1380

Thr Gln Ala Ala Ala Ile
1385

<210> 80
<211> 148
<212> PRT
<213> Mouse

<400> 80

Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
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Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Phe
20 25 30

Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg
35 40 45

Glu Asp Ser Pro Ala Ala Lys Ser Leu Ser Leu Gln Glu Gly Asp Gln
50 55 60

Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala
65 70 75 80

Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu
85 90 95

Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val
100 105 110

Ser Gly Tyr Glu Met Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Val
115 120 125

Arg Val Leu Ser Pro Val Pro Val Gln Asp Ser Pro Ser Asp Arg Val
130 135 140

Ala Ala Ala Pro

145

<210> 81
<211> 377
<212> PRT
<213> Caenorhabditiselegans

<400> 81

Met Ser Val Phe Arg Phe Leu Leu Phe Leu Ser Leu Leu Val Gly Ser
1 5 10 15

Asn Ala Phe Val Lys Pro Gln Tyr Asn Val Thr Gly Gln Ile Asp Ser
20 25 30

Ala Leu Gln Arg Phe Phe Gly Ile Thr Leu Pro Ser Leu Lys Ile Pro
35 40 45

Asp Leu Leu Asn Pro Asp Lys Lys Arg Asn Pro Pro Ser Val Gly Gln
50 55 60

Leu Lys Lys Thr Ser Phe Pro Leu Cys Asn Val Asn Leu Pro Pro Ile
65 70 75 80

Phe Phe Thr Ile Ser Leu Phe Arg Ile Lys Leu Pro Asn Leu Ile Pro
85 90 95

Thr Ala Leu Pro Val Ile Lys Leu Pro Thr Ile Lys Ile Pro Asn Ile
100 105 110

Leu Pro Thr Leu Pro Thr Ile Lys Val Pro Thr Ile Lys Ile Pro Asp
115 120 125

Ile Ile Pro Ile Thr Leu Pro Thr Ile Lys Ile Pro Glu Val Val Pro
130 135 140

Thr Asn Leu Pro Thr Val Glu Ile Pro His Phe Ile Pro Lys Thr Leu
145 150 155 160

Pro Thr Val Lys Ile Pro Asn Ile Ile Pro Thr Asn Phe Pro Thr Ile
165 170 175

Glu Thr Pro Asp Ile Ile Pro Lys Ile Leu Pro Thr Ile Lys Ile Pro
180 185 190

Glu Ile Ile Pro Leu Thr Leu Pro Thr Val Lys Ile Pro Asp Ile Ile
195 200 205

25167341

Pro Ile Thr Leu Pro Thr Ile Lys Ile Pro Glu Ile Val Pro Thr Lys
 210 215 220

Leu Pro Thr Val Glu Val Pro Asp Thr Ile Pro Lys Thr Leu Pro Thr
 225 230 235 240

Thr Lys Ile Pro Asp Ile Val Pro Ile Thr Ser Pro Thr Val Lys Ile
 245 250 255

Pro Gln Ile Ile Pro Thr Ile Lys Ile Pro Asp Ile Ile Pro Lys Asn
 260 265 270

Leu Ser Thr Leu Gly Pro Ile Lys Leu Pro Thr Ile Lys Leu Pro Thr
 275 280 285

Gly Asn Met Val Cys Asp Ile Cys Glu Lys Val Ile Gly Val Leu Thr
 290 295 300

Thr Arg Leu Leu Glu Ile Ile Gln Lys Phe Arg Val Glu Ala Asp Lys
 305 310 315 320

Phe Leu Thr Lys Leu Cys Thr Ser Leu Thr Ser Asn Pro Lys Thr Leu
 325 330 335

Thr Val Gly Thr Met Cys Val Met Phe Lys Gly Asn Ile Met Asp Thr
 340 345 350

Ile Phe Lys Gly Phe Asp Gly Leu Lys Lys Asn Leu Glu Pro Val Ser
 355 360 365

Phe Cys Lys His Val Pro Phe Cys Lys
 370 375

<210> 82
 <211> 1383
 <212> PRT
 <213> Rat

<400> 82

Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
 1 5 10 15

Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Phe
 20 25 30

Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg

| | | | | | | | | | | | | |
|---|-----|-----|--|--|--|--|--|--|--|--|--|--|
| 35 | 40 | 45 | | | | | | | | | | |
| Glu Asp Ser Pro Ala Ala Lys Ser Leu Ser Leu Gln Glu Gly Asp Gln | | | | | | | | | | | | |
| 50 | 55 | 60 | | | | | | | | | | |
| Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala | | | | | | | | | | | | |
| 65 | 70 | 75 | | | | | | | | | | |
| Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu | | | | | | | | | | | | |
| | 85 | 90 | | | | | | | | | | |
| Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val | | | | | | | | | | | | |
| | 100 | 105 | | | | | | | | | | |
| Ser Gly Tyr Glu Met Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Asn | | | | | | | | | | | | |
| | 115 | 120 | | | | | | | | | | |
| Ile Gln Ser Leu Ser Pro Val Lys Lys Lys Lys Met Val Ile Gly Thr | | | | | | | | | | | | |
| | 130 | 135 | | | | | | | | | | |
| Leu Gly Thr Pro Ala Asp Leu Ala Pro Val Asp Val Glu Phe Ser Phe | | | | | | | | | | | | |
| 145 | 150 | 155 | | | | | | | | | | |
| Pro Lys Phe Ser Arg Leu Arg Arg Gly Leu Lys Ala Asp Ala Val Lys | | | | | | | | | | | | |
| | 165 | 170 | | | | | | | | | | |
| Gly Pro Val Pro Ala Ala Pro Ala Arg Arg Arg Leu Gln Leu Pro Arg | | | | | | | | | | | | |
| | 180 | 185 | | | | | | | | | | |
| Leu Arg Val Arg Glu Val Ala Glu Glu Ala Gln Val Ala Arg Met Ala | | | | | | | | | | | | |
| | 195 | 200 | | | | | | | | | | |
| Ala Ala Ala Pro Pro Ser Arg Lys Ala Lys Ser Glu Ala Glu Val Ala | | | | | | | | | | | | |
| | 210 | 215 | | | | | | | | | | |
| Thr Gly Ala Gly Phe Thr Ala Pro Gln Ile Glu Leu Val Gly Pro Arg | | | | | | | | | | | | |
| 225 | 230 | 235 | | | | | | | | | | |
| Leu Pro Ser Ala Glu Val Gly Val Pro Lys Val Ser Val Pro Lys Gly | | | | | | | | | | | | |
| | 245 | 250 | | | | | | | | | | |
| Thr Pro Ser Thr Glu Ala Ala Ser Gly Phe Ala Leu His Leu Pro Thr | | | | | | | | | | | | |
| | 260 | 265 | | | | | | | | | | |
| Leu Gly Leu Gly Ala Pro Ala Ala Pro Ala Val Glu Pro Pro Thr Thr | | | | | | | | | | | | |
| | 275 | 280 | | | | | | | | | | |
| | | 285 | | | | | | | | | | |

Gly Ile Gln Val Pro Gln Val Glu Leu Pro Thr Leu Pro Ser Leu Pro
 290 295 300

Thr Leu Pro Thr Leu Pro Cys Leu Asp Thr Gln Glu Gly Ala Ala Val
 305 310 315 320

Val Lys Val Pro Thr Leu Asp Val Ala Ala Pro Ser Val Glu Val Asp
 325 330 335

Leu Ala Leu Pro Gly Ala Glu Val Glu Ala Gln Gly Glu Val Pro Glu
 340 345 350

Val Ala Leu Lys Met Pro Arg Leu Ser Phe Pro Arg Phe Gly Val Arg
 355 360 365

Gly Lys Glu Ala Thr Glu Ala Lys Val Val Lys Gly Ser Pro Glu Ala
 370 375 380

Lys Ala Lys Gly Pro Arg Leu Arg Met Pro Thr Phe Gly Leu Ser Leu
 385 390 395 400

Leu Glu Ser Arg Pro Ser Gly Pro Glu Val Ala Ala Glu Ser Lys Leu
 405 410 415

Lys Leu Pro Thr Leu Lys Met Pro Ser Phe Gly Ile Ser Val Ala Gly
 420 425 430

Pro Glu Val Lys Ala Pro Lys Gly Pro Glu Val Lys Leu Pro Lys Val
 435 440 445

Pro Glu Ile Lys Leu Pro Lys Ala Pro Glu Ala Ala Ile Pro Asp Val
 450 455 460

Gln Leu Pro Glu Val Gln Leu Pro Lys Met Ser Asp Met Lys Leu Pro
 465 470 475 480

Lys Ile Pro Glu Met Ala Val Pro Asp Val His Leu Pro Glu Val Lys
 485 490 495

Leu Pro Lys Val Pro Glu Met Lys Val Pro Glu Met Lys Leu Pro Lys
 500 505 510

Ile Pro Glu Met Ala Val Pro Asp Val His Leu Pro Asp Ile Gln Leu
 515 520 525

Pro Lys Val Pro Glu Met Lys Leu Pro Asp Met Lys Leu Pro Lys Val
 530 535 540

Pro Glu Met Ala Val Pro Asp Val His Leu Pro Asp Ile Gln Leu Pro
 545 550 555 560

Lys Val Pro Glu Met Lys Leu Pro Asp Met Lys Leu Pro Lys Val Pro
 565 570 575

Glu Met Ala Val Pro Asp Val Arg Ile Pro Glu Val Gln Leu Pro Lys
 580 585 590

Val Ser Glu Val Lys Leu Pro Lys Ile Pro Asp Met Ala Val Pro Asp
 595 600 605

Val Arg Leu Pro Glu Leu Gln Leu Pro Lys Met Ser Glu Val Lys Leu
 610 615 620

Pro Lys Ile Pro Asp Met Ala Val Pro Asp Val Arg Leu Pro Glu Val
 625 630 635 640

Gln Leu Pro Lys Val Ser Glu Leu Lys Leu Pro Lys Val Pro Glu Met
 645 650 655

Thr Met Pro Asp Ile Arg Leu Pro Glu Val Gln Leu Pro Lys Val Pro
 660 665 670

Asp Ile Lys Leu Pro Glu Ile Lys Leu Pro Lys Val Pro Glu Met Ala
 675 680 685

Val Pro Asp Val Pro Leu Pro Glu Leu Gln Leu Pro Lys Val Pro Gln
 690 695 700

Val Pro Asp Val His Leu Pro Lys Val Pro Glu Met Lys Leu Pro Lys
 705 710 715 720

Val Pro Glu Ala Gln Arg Lys Ser Ala Gly Ala Glu Gln Ala Glu Lys
 725 730 735

Thr Glu Phe Ser Phe Lys Leu Pro Lys Met Thr Val Pro Lys Leu Gly
 740 745 750

Lys Val Thr Lys Pro Gly Glu Ala Gly Ile Glu Val Pro Asp Lys Leu
 755 760 765

Leu Ile Leu Pro Cys Leu Gln Pro Glu Val Gly Thr Glu Val Ala Arg
 770 775 780

Val Gly Val Pro Ser Leu Ser Leu Pro Ser Val Glu Leu Asp Leu Pro
 785 790 795 800

Gly Ala Leu Gly Leu Glu Gly Gln Val Gln Glu Ala Val Ser Gly Lys
 805 810 815

Val Glu Lys Pro Glu Gly Pro Arg Val Ala Val Gly Thr Gly Glu Ala
 820 825 830

Gly Phe Arg Val Pro Ser Val Glu Ile Val Asn Pro Gln Leu Pro Thr
 835 840 845

Val Glu Val Lys Lys Glu Gln Leu Glu Met Val Glu Met Lys Val Lys
 850 855 860

Pro Thr Ser Lys Phe Ser Leu Pro Lys Phe Gly Leu Ser Gly Pro Lys
 865 870 875 880

Ala Val Lys Ala Glu Val Glu Gly Pro Gly Arg Ala Thr Lys Leu Lys
 885 890 895

Val Ser Lys Phe Ala Ile Ser Leu Pro Arg Ala Arg Ala Gly Thr Asp
 900 905 910

Ala Asp Ala Lys Gly Ala Gly Glu Ala Gly Leu Leu Pro Ala Leu Asp
 915 920 925

Leu Ser Ile Pro Gln Leu Ser Leu Asp Ala Gln Leu Pro Ser Gly Lys
 930 935 940

Val Glu Val Ala Gly Ala Glu Ser Lys Pro Lys Gly Ser Arg Phe Ala
 945 950 955 960

Leu Pro Lys Phe Gly Ala Lys Gly Arg Asp Ser Glu Ala Asp Val Leu
 965 970 975

Val Ala Gly Glu Ala Glu Leu Glu Gly Lys Gly Trp Gly Trp Asp Gly
 980 985 990

Lys Val Lys Met Pro Lys Leu Lys Met Pro Ser Phe Gly Leu Ser Arg
 995 1000 1005

Gly Lys Glu Ala Glu Ile Gln Asp Gly Arg Val Ser Pro Gly Glu

| | | | | |
|---|--|------|--|------|
| 1010 | | 1015 | | 1020 |
| Lys Leu Glu Ala Ile Ala Gly Gln Leu Lys Ile Pro Glu Val Glu | | | | |
| 1025 | | 1030 | | 1035 |
| Leu Val Thr Pro Gly Ala Gln Glu Thr Glu Lys Val Thr Ser Gly | | | | |
| 1040 | | 1045 | | 1050 |
| Val Lys Pro Ser Gly Leu Gln Val Ser Thr Thr Arg Gln Val Val | | | | |
| 1055 | | 1060 | | 1065 |
| Ala Glu Gly Gln Glu Gly Ala Gln Arg Val Ser Ser Leu Gly Ile | | | | |
| 1070 | | 1075 | | 1080 |
| Ser Leu Pro Gln Val Glu Leu Ala Ser Phe Gly Glu Ala Gly Pro | | | | |
| 1085 | | 1090 | | 1095 |
| Glu Ile Ala Ala Pro Ser Ala Glu Gly Thr Val Gly Ser Arg Ile | | | | |
| 1100 | | 1105 | | 1110 |
| Gln Val Pro Gln Val Met Leu Glu Leu Pro Gly Thr Gln Val Ala | | | | |
| 1115 | | 1120 | | 1125 |
| Gly Gly Asp Leu Leu Val Gly Glu Gly Ile Phe Lys Met Pro Thr | | | | |
| 1130 | | 1135 | | 1140 |
| Val Thr Val Pro Gln Leu Glu Leu Asp Val Gly Leu Gly His Glu | | | | |
| 1145 | | 1150 | | 1155 |
| Ala Gln Ala Gly Glu Thr Ala Lys Ser Glu Gly Gly Leu Lys Leu | | | | |
| 1160 | | 1165 | | 1170 |
| Lys Leu Pro Thr Leu Gly Ala Gly Gly Lys Gly Glu Gly Ala Glu | | | | |
| 1175 | | 1180 | | 1185 |
| Ala Gln Ser Pro Glu Ala Gln His Thr Phe His Ile Ser Leu Pro | | | | |
| 1190 | | 1195 | | 1200 |
| Asp Val Glu Leu Thr Ser Pro Val Ser Ser His Ala Glu Tyr Gln | | | | |
| 1205 | | 1210 | | 1215 |
| Val Val Glu Gly Asp Gly Asp Gly Gly His Lys Leu Lys Val Arg | | | | |
| 1220 | | 1225 | | 1230 |
| Leu Pro Leu Phe Gly Leu Ala Arg Ala Lys Glu Gly Ile Glu Thr | | | | |
| 1235 | | 1240 | | 1245 |

Gly Glu Lys Val Lys Ser Pro Lys Leu Arg Leu Pro Arg Val Gly
1250 1255 1260

Phe Ser Gln Ser Glu Ser Ala Ser Gly Glu Gly Ser Pro Ser Pro
1265 1270 1275

Glu Glu Glu Glu Glu Gly Ser Gly Glu Gly Ala Ser Gly Arg Arg
1280 1285 1290

Gly Arg Val Arg Val Arg Leu Pro Arg Val Gly Leu Ala Ser Pro
1295 1300 1305

Ser Lys Gly Ser Lys Gly Gln Glu Gly Asp Ala Ala Ser Lys Ser
1310 1315 1320

Pro Val Gly Glu Lys Ser Pro Lys Phe Arg Phe Pro Arg Val Ser
1325 1330 1335

Leu Ser Pro Lys Ala Arg Ser Gly Ser Lys Asp Arg Glu Glu Gly
1340 1345 1350

Gly Phe Arg Val Arg Leu Pro Ser Val Gly Phe Ser Glu Thr Ala
1355 1360 1365

Ala Pro Gly Ser Ala Arg Ile Glu Gly Thr Gln Ala Ala Ala Ile
1370 1375 1380

<210> 83
<211> 370
<212> PRT
<213> Arabidopsisthaliana

<400> 83

Met Ala Leu Met Lys Lys Ser Leu Ser Ala Ala Leu Leu Ser Ser Pro
1 5 10 15

Leu Leu Ile Ile Cys Leu Ile Ala Leu Leu Ala Asp Pro Phe Ser Val
20 25 30

Gly Ala Arg Arg Leu Leu Glu Asp Pro Lys Pro Glu Ile Pro Lys Leu
35 40 45

Pro Glu Leu Pro Lys Phe Glu Val Pro Lys Leu Pro Glu Phe Pro Lys
50 55 60

Pro Glu Leu Pro Lys Leu Pro Glu Phe Pro Lys Pro Glu Leu Pro Lys
65 70 75 80

Ile Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Val Pro Glu Ile Pro
85 90 95

Lys Pro Glu Glu Thr Lys Leu Pro Asp Ile Pro Lys Leu Glu Leu Pro
100 105 110

Lys Phe Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Met Pro Glu Ile
115 120 125

Pro Lys Pro Glu Leu Pro Lys Val Pro Glu Ile Gln Lys Pro Glu Leu
130 135 140

Pro Lys Met Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Phe Pro Glu
145 150 155 160

Ile Pro Lys Pro Asp Leu Pro Lys Phe Pro Glu Asn Ser Lys Pro Glu
165 170 175

Val Pro Lys Leu Met Glu Thr Glu Lys Pro Glu Ala Pro Lys Val Pro
180 185 190

Glu Ile Pro Lys Pro Glu Leu Pro Lys Leu Pro Glu Val Pro Lys Leu
195 200 205

Glu Ala Pro Lys Val Pro Glu Ile Gln Lys Pro Glu Leu Pro Lys Met
210 215 220

Pro Glu Leu Pro Lys Met Pro Glu Ile Gln Lys Pro Glu Leu Pro Lys
225 230 235 240

Leu Pro Glu Val Pro Lys Leu Glu Ala Pro Lys Val Pro Glu Ile Gln
245 250 255

Lys Pro Glu Leu Pro Lys Met Pro Glu Leu Pro Lys Met Pro Glu Ile
260 265 270

Gln Lys Pro Glu Leu Pro Lys Met Pro Glu Ile Gln Lys Pro Glu Leu
275 280 285

Pro Lys Val Pro Glu Val Pro Lys Pro Glu Leu Pro Thr Val Pro Glu
290 295 300

Val Pro Lys Ser Glu Ala Pro Lys Phe Pro Glu Ile Pro Lys Pro Glu

305 310 315 320
 Leu Pro Lys Ile Pro Glu Val Pro Lys Pro Glu Leu Pro Lys Val Pro
 325 330 335
 Glu Ile Thr Lys Pro Ala Val Pro Glu Ile Pro Lys Pro Glu Leu Pro
 340 345 350
 Thr Met Pro Gln Leu Pro Lys Leu Pro Glu Phe Pro Lys Val Pro Gly
 355 360 365
 Thr Pro
 370

 <210> 84
 <211> 148
 <212> PRT
 <213> Mouse

 <400> 84
 Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
 1 5 10 15
 Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Phe
 20 25 30
 Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg
 35 40 45
 Glu Asp Ser Pro Ala Ala Lys Ser Leu Ser Leu Gln Glu Gly Asp Gln
 50 55 60
 Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala
 65 70 75 80
 Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu
 85 90 95
 Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val
 100 105 110
 Ser Gly Tyr Glu Met Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Val
 115 120 125
 Arg Val Leu Ser Pro Val Pro Val Gln Asp Ser Pro Ser Asp Arg Val
 130 135 140

Ala Ala Ala Pro
145

<210> 85
<211> 1391
<212> PRT
<213> Mouse

<400> 85

Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
1 5 10 15

Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Phe
20 25 30

Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg
35 40 45

Glu Asp Ser Pro Ala Ala Lys Ser Leu Ser Leu Gln Glu Gly Asp Gln
50 55 60

Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala
65 70 75 80

Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu
85 90 95

Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val
100 105 110

Ser Gly Tyr Glu Met Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Asn
115 120 125

Ile Gln Ser Leu Ala Pro Val Lys Lys Lys Lys Met Val Thr Gly Ala
130 135 140

Leu Gly Thr Pro Ala Asp Leu Ala Pro Val Asp Val Glu Phe Ser Phe
145 150 155 160

Pro Lys Phe Ser Arg Leu Arg Arg Gly Leu Lys Ala Glu Ala Val Lys
165 170 175

Gly Pro Val Pro Ala Ala Pro Ala Arg Arg Arg Leu Gln Leu Pro Arg
180 185 190

Leu Arg Val Arg Glu Val Ala Glu Glu Ala Gln Val Ala Arg Met Ala

| | | | | |
|---|-----|-----|-----|-----|
| 195 | | 200 | | 205 |
| Ala Ala Ala Pro Pro Pro Arg Lys Ala Lys Ala Glu Ala Glu Ala Ala | 210 | 215 | 220 | |
| Thr Gly Ala Gly Phe Thr Ala Pro Gln Ile Glu Leu Val Gly Pro Arg | 225 | 230 | 235 | 240 |
| Leu Pro Ser Ala Glu Val Gly Val Pro Gln Val Ser Val Pro Lys Gly | 245 | 250 | 255 | |
| Thr Pro Ser Thr Glu Ala Ala Ser Gly Phe Ala Leu His Leu Pro Thr | 260 | 265 | 270 | |
| Leu Gly Leu Gly Ala Pro Ala Ala Pro Ala Val Glu Pro Pro Ala Thr | 275 | 280 | 285 | |
| Gly Ile Gln Val Pro Gln Val Glu Leu Pro Thr Leu Pro Ser Leu Pro | 290 | 295 | 300 | |
| Thr Leu Pro Thr Leu Pro Cys Leu Asp Thr Gln Glu Gly Ala Ala Val | 305 | 310 | 315 | 320 |
| Val Lys Val Pro Thr Leu Asp Val Ala Ala Pro Ser Met Gly Val Asp | 325 | 330 | 335 | |
| Leu Ala Leu Pro Gly Ala Glu Val Glu Ala Gln Gly Glu Val Pro Glu | 340 | 345 | 350 | |
| Val Ala Leu Lys Met Pro Arg Leu Ser Phe Pro Arg Phe Gly Ile Arg | 355 | 360 | 365 | |
| Gly Lys Glu Ala Thr Glu Ala Lys Val Val Lys Gly Ser Pro Glu Ala | 370 | 375 | 380 | |
| Lys Ala Lys Gly Pro Arg Leu Arg Met Pro Thr Phe Gly Leu Ser Leu | 385 | 390 | 395 | 400 |
| Leu Glu Pro Arg Pro Ser Gly Pro Glu Ala Val Ala Glu Ser Lys Leu | 405 | 410 | 415 | |
| Lys Leu Pro Thr Leu Lys Met Pro Ser Phe Gly Ile Gly Val Ala Gly | 420 | 425 | 430 | |
| Pro Glu Val Lys Ala Pro Thr Gly Pro Glu Val Lys Leu Pro Lys Val | 435 | 440 | 445 | |

Pro Glu Val Lys Leu Pro Lys Val Pro Glu Ala Ala Ile Pro Asp Val
 450 455 460

Gln Leu Pro Glu Val Gln Leu Pro Lys Met Ser Asp Met Lys Leu Pro
 465 470 475 480

Lys Ile Pro Glu Met Val Val Pro Asp Val Arg Leu Pro Glu Val Gln
 485 490 495

Leu Pro Lys Val Pro Glu Met Lys Val Pro Glu Met Lys Leu Pro Lys
 500 505 510

Trp Pro Glu Met Ala Val Pro Asp Val His Leu Pro Asp Val Gln Leu
 515 520 525

Pro Lys Val Pro Glu Met Lys Leu Pro Lys Val Pro Glu Met Ala Val
 530 535 540

Pro Asp Val His Leu Pro Asp Val Gln Leu Pro Lys Val Pro Glu Met
 545 550 555 560

Lys Leu Pro Glu Met Lys Leu Pro Lys Val Pro Glu Met Ala Val Pro
 565 570 575

Asp Val Arg Leu Pro Glu Val Gln Leu Pro Lys Val Ser Glu Val Lys
 580 585 590

Leu Pro Lys Met Pro Glu Met Ala Val Pro Asp Val His Leu Pro Glu
 595 600 605

Leu Gln Leu Pro Lys Met Ser Glu Val Lys Leu Pro Lys Met Pro Glu
 610 615 620

Met Ala Val Pro Asp Val Arg Leu Pro Glu Val Gln Leu Pro Lys Val
 625 630 635 640

Ser Glu Met Lys Leu Pro Lys Met Pro Glu Met Thr Met Pro Asp Ile
 645 650 655

Arg Leu Pro Glu Val Gln Leu Pro Lys Val Pro Asp Ile Lys Leu Pro
 660 665 670

Glu Met Lys Leu Pro Glu Ile Lys Leu Pro Lys Val Pro Asp Met Ala
 675 680 685

Val Pro Asp Val Pro Leu Pro Glu Leu Gln Leu Pro Lys Val Ser Asp
690 695 700
Ile Arg Leu Pro Glu Met Gln Val Ser Gln Val Pro Glu Val Gln Leu
705 710 715 720
Pro Lys Met Pro Glu Met Lys Leu Ser Lys Val Pro Glu Val Gln Arg
725 730 735
Lys Ser Ala Gly Ala Glu Gln Ala Lys Gly Thr Glu Phe Ser Phe Lys
740 745 750
Leu Pro Lys Met Thr Met Pro Lys Leu Gly Lys Val Gly Lys Pro Gly
755 760 765
Glu Ala Ser Ile Glu Val Pro Asp Lys Leu Met Thr Leu Pro Cys Leu
770 775 780
Gln Pro Glu Val Gly Thr Glu Ala Ser His Val Gly Val Pro Ser Leu
785 790 795 800
Ser Leu Pro Ser Val Glu Leu Asp Leu Pro Gly Ala Leu Gly Leu Glu
805 810 815
Gly Gln Val Gln Glu Ala Val Pro Gly Lys Val Glu Lys Pro Glu Gly
820 825 830
Pro Arg Val Ala Val Gly Val Gly Glu Val Gly Phe Arg Val Pro Ser
835 840 845
Val Glu Ile Val Thr Pro Gln Leu Pro Thr Val Glu Val Glu Lys Glu
850 855 860
Gln Leu Glu Met Val Glu Met Lys Val Lys Pro Ser Ser Lys Phe Ser
865 870 875 880
Leu Pro Lys Phe Gly Leu Ser Gly Pro Lys Ala Val Lys Gly Glu Val
885 890 895
Glu Gly Pro Gly Arg Ala Thr Lys Leu Lys Val Ser Lys Phe Thr Ile
900 905 910
Ser Leu Pro Lys Ala Arg Ala Gly Thr Glu Ala Glu Ala Lys Gly Ala
915 920 925

Gly Glu Ala Gly Leu Leu Pro Ala Leu Asp Leu Ser Ile Pro Gln Leu
 930 935 940

Ser Leu Asp Ala Gln Leu Pro Ser Gly Lys Val Glu Val Ala Asp Ser
 945 950 955 960

Lys Pro Lys Ser Ser Arg Phe Ala Leu Pro Lys Phe Gly Val Lys Gly
 965 970 975

Arg Asp Ser Glu Ala Asp Val Leu Val Ala Gly Glu Ala Glu Leu Glu
 980 985 990

Gly Lys Gly Trp Gly Trp Asp Gly Lys Val Lys Met Pro Lys Leu Lys
 995 1000 1005

Met Pro Ser Phe Gly Leu Ser Arg Gly Lys Glu Ala Glu Thr Gln
 1010 1015 1020

Asp Gly Arg Val Ser Pro Gly Glu Lys Leu Glu Ala Ile Ala Gly
 1025 1030 1035

Gln Leu Lys Ile Pro Ala Val Glu Leu Val Thr Pro Gly Ala Gln
 1040 1045 1050

Glu Thr Glu Lys Val Thr Ser Gly Val Lys Pro Ser Gly Leu Gln
 1055 1060 1065

Val Ser Thr Thr Gly Gln Val Val Ala Glu Gly Gln Glu Ser Val
 1070 1075 1080

Gln Arg Val Ser Thr Leu Gly Ile Ser Leu Pro Gln Val Glu Leu
 1085 1090 1095

Ala Ser Phe Gly Glu Ala Gly Pro Glu Ile Val Ala Pro Ser Ala
 1100 1105 1110

Glu Gly Thr Ala Gly Ser Arg Val Gln Val Pro Gln Val Met Leu
 1115 1120 1125

Glu Leu Pro Gly Thr Gln Val Ala Gly Gly Asp Leu Leu Val Gly
 1130 1135 1140

Glu Gly Ile Phe Lys Met Pro Thr Val Thr Val Pro Gln Leu Glu
 1145 1150 1155

Leu Asp Val Gly Leu Gly His Glu Ala Gln Ala Gly Glu Ala Ala

| | | | | |
|-------------------------|---------------------|-----------------|--|------|
| 1160 | | 1165 | | 1170 |
| Lys Ser Glu Gly Gly Ile | Lys Leu Lys Leu Pro | Thr Leu Gly Thr | | |
| 1175 | 1180 | 1185 | | |
| Gly Ser Arg Gly Glu Gly | Val Glu Pro Gln Gly | Pro Glu Ala Gln | | |
| 1190 | 1195 | 1200 | | |
| Arg Thr Phe His Leu Ser | Leu Pro Asp Val Glu | Leu Thr Ser Pro | | |
| 1205 | 1210 | 1215 | | |
| Val Ser Ser His Ala Glu | Tyr Gln Val Val Glu | Gly Asp Gly Asp | | |
| 1220 | 1225 | 1230 | | |
| Gly Gly His Lys Leu Lys | Val Arg Leu Pro Leu | Phe Gly Leu Ala | | |
| 1235 | 1240 | 1245 | | |
| Lys Ala Lys Glu Gly Ile | Glu Val Gly Glu Lys | Val Lys Ser Pro | | |
| 1250 | 1255 | 1260 | | |
| Lys Leu Arg Leu Pro Arg | Val Gly Phe Ser Gln | Ser Glu Ser Val | | |
| 1265 | 1270 | 1275 | | |
| Ser Gly Glu Gly Ser Pro | Ser Pro Glu Glu Glu | Glu Glu Gly Ser | | |
| 1280 | 1285 | 1290 | | |
| Gly Glu Gly Ala Ser Ser | Arg Arg Gly Arg Val | Arg Val Arg Leu | | |
| 1295 | 1300 | 1305 | | |
| Pro Arg Val Gly Leu Ala | Ser Pro Ser Lys Val | Ser Lys Gly Gln | | |
| 1310 | 1315 | 1320 | | |
| Glu Gly Asp Ala Thr Ser | Lys Ser Pro Val Gly | Glu Lys Ser Pro | | |
| 1325 | 1330 | 1335 | | |
| Lys Phe Arg Phe Pro Arg | Val Ser Leu Ser Pro | Lys Ala Arg Ser | | |
| 1340 | 1345 | 1350 | | |
| Gly Ser Arg Asp Arg Glu | Glu Gly Gly Phe Arg | Val Arg Leu Pro | | |
| 1355 | 1360 | 1365 | | |
| Ser Val Gly Phe Ser Glu | Thr Ala Val Pro Gly | Ser Thr Arg Ile | | |
| 1370 | 1375 | 1380 | | |
| Glu Gly Thr Gln Ala Ala | Ala Ile | | | |
| 1385 | 1390 | | | |

<210> 86
 <211> 1383
 <212> PRT
 <213> Rat

<400> 86

Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
 1 5 10 15

Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Phe
 20 25 30

Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg
 35 40 45

Glu Asp Ser Pro Ala Ala Lys Ser Leu Ser Leu Gln Glu Gly Asp Gln
 50 55 60

Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala
 65 70 75 80

Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu
 85 90 95

Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val
 100 105 110

Ser Gly Tyr Glu Met Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Asn
 115 120 125

Ile Gln Ser Leu Ser Pro Val Lys Lys Lys Lys Met Val Ile Gly Thr
 130 135 140

Leu Gly Thr Pro Ala Asp Leu Ala Pro Val Asp Val Glu Phe Ser Phe
 145 150 155 160

Pro Lys Phe Ser Arg Leu Arg Arg Gly Leu Lys Ala Asp Ala Val Lys
 165 170 175

Gly Pro Val Pro Ala Ala Pro Ala Arg Arg Arg Leu Gln Leu Pro Arg
 180 185 190

Leu Arg Val Arg Glu Val Ala Glu Glu Ala Gln Val Ala Arg Met Ala
 195 200 205

Ala Ala Ala Pro Pro Ser Arg Lys Ala Lys Ser Glu Ala Glu Val Ala
 210 215 220
 Thr Gly Ala Gly Phe Thr Ala Pro Gln Ile Glu Leu Val Gly Pro Arg
 225 230 235 240
 Leu Pro Ser Ala Glu Val Gly Val Pro Lys Val Ser Val Pro Lys Gly
 245 250 255
 Thr Pro Ser Thr Glu Ala Ala Ser Gly Phe Ala Leu His Leu Pro Thr
 260 265 270
 Leu Gly Leu Gly Ala Pro Ala Ala Pro Ala Val Glu Pro Pro Thr Thr
 275 280 285
 Gly Ile Gln Val Pro Gln Val Glu Leu Pro Thr Leu Pro Ser Leu Pro
 290 295 300
 Thr Leu Pro Thr Leu Pro Cys Leu Asp Thr Gln Glu Gly Ala Ala Val
 305 310 315 320
 Val Lys Val Pro Thr Leu Asp Val Ala Ala Pro Ser Val Glu Val Asp
 325 330 335
 Leu Ala Leu Pro Gly Ala Glu Val Glu Ala Gln Gly Glu Val Pro Glu
 340 345 350
 Val Ala Leu Lys Met Pro Arg Leu Ser Phe Pro Arg Phe Gly Val Arg
 355 360 365
 Gly Lys Glu Ala Thr Glu Ala Lys Val Val Lys Gly Ser Pro Glu Ala
 370 375 380
 Lys Ala Lys Gly Pro Arg Leu Arg Met Pro Thr Phe Gly Leu Ser Leu
 385 390 395 400
 Leu Glu Ser Arg Pro Ser Gly Pro Glu Val Ala Ala Glu Ser Lys Leu
 405 410 415
 Lys Leu Pro Thr Leu Lys Met Pro Ser Phe Gly Ile Ser Val Ala Gly
 420 425 430
 Pro Glu Val Lys Ala Pro Lys Gly Pro Glu Val Lys Leu Pro Lys Val
 435 440 445
 Pro Glu Ile Lys Leu Pro Lys Ala Pro Glu Ala Ala Ile Pro Asp Val

| | | | | |
|---|---|-----|-----|-----|
| 450 | | 455 | | 460 |
| Gln Leu Pro Glu Val | Gln Leu Pro Lys Met Ser Asp Met Lys Leu Pro | | | |
| 465 | 470 | 475 | | 480 |
| Lys Ile Pro Glu Met Ala Val Pro Asp Val His Leu Pro Glu Val Lys | | | | |
| 485 | 490 | | | 495 |
| Leu Pro Lys Val Pro Glu Met Lys Val Pro Glu Met Lys Leu Pro Lys | | | | |
| 500 | 505 | | | 510 |
| Ile Pro Glu Met Ala Val Pro Asp Val His Leu Pro Asp Ile Gln Leu | | | | |
| 515 | 520 | | 525 | |
| Pro Lys Val Pro Glu Met Lys Leu Pro Asp Met Lys Leu Pro Lys Val | | | | |
| 530 | 535 | | 540 | |
| Pro Glu Met Ala Val Pro Asp Val His Leu Pro Asp Ile Gln Leu Pro | | | | |
| 545 | 550 | | 555 | 560 |
| Lys Val Pro Glu Met Lys Leu Pro Asp Met Lys Leu Pro Lys Val Pro | | | | |
| 565 | 570 | | | 575 |
| Glu Met Ala Val Pro Asp Val Arg Ile Pro Glu Val Gln Leu Pro Lys | | | | |
| 580 | 585 | | | 590 |
| Val Ser Glu Val Lys Leu Pro Lys Ile Pro Asp Met Ala Val Pro Asp | | | | |
| 595 | 600 | | 605 | |
| Val Arg Leu Pro Glu Leu Gln Leu Pro Lys Met Ser Glu Val Lys Leu | | | | |
| 610 | 615 | | 620 | |
| Pro Lys Ile Pro Asp Met Ala Val Pro Asp Val Arg Leu Pro Glu Val | | | | |
| 625 | 630 | | 635 | 640 |
| Gln Leu Pro Lys Val Ser Glu Leu Lys Leu Pro Lys Val Pro Glu Met | | | | |
| 645 | 650 | | | 655 |
| Thr Met Pro Asp Ile Arg Leu Pro Glu Val Gln Leu Pro Lys Val Pro | | | | |
| 660 | 665 | | | 670 |
| Asp Ile Lys Leu Pro Glu Ile Lys Leu Pro Lys Val Pro Glu Met Ala | | | | |
| 675 | 680 | | 685 | |
| Val Pro Asp Val Pro Leu Pro Glu Leu Gln Leu Pro Lys Val Pro Gln | | | | |
| 690 | 695 | | 700 | |

Val Pro Asp Val His Leu Pro Lys Val Pro Glu Met Lys Leu Pro Lys
705 710 715 720

Val Pro Glu Ala Gln Arg Lys Ser Ala Gly Ala Glu Gln Ala Glu Lys
725 730 735

Thr Glu Phe Ser Phe Lys Leu Pro Lys Met Thr Val Pro Lys Leu Gly
740 745 750

Lys Val Thr Lys Pro Gly Glu Ala Gly Ile Glu Val Pro Asp Lys Leu
755 760 765

Leu Ile Leu Pro Cys Leu Gln Pro Glu Val Gly Thr Glu Val Ala Arg
770 775 780

Val Gly Val Pro Ser Leu Ser Leu Pro Ser Val Glu Leu Asp Leu Pro
785 790 795 800

Gly Ala Leu Gly Leu Glu Gly Gln Val Gln Glu Ala Val Ser Gly Lys
805 810 815

Val Glu Lys Pro Glu Gly Pro Arg Val Ala Val Gly Thr Gly Glu Ala
820 825 830

Gly Phe Arg Val Pro Ser Val Glu Ile Val Asn Pro Gln Leu Pro Thr
835 840 845

Val Glu Val Lys Lys Glu Gln Leu Glu Met Val Glu Met Lys Val Lys
850 855 860

Pro Thr Ser Lys Phe Ser Leu Pro Lys Phe Gly Leu Ser Gly Pro Lys
865 870 875 880

Ala Val Lys Ala Glu Val Glu Gly Pro Gly Arg Ala Thr Lys Leu Lys
885 890 895

Val Ser Lys Phe Ala Ile Ser Leu Pro Arg Ala Arg Ala Gly Thr Asp
900 905 910

Ala Asp Ala Lys Gly Ala Gly Glu Ala Gly Leu Leu Pro Ala Leu Asp
915 920 925

Leu Ser Ile Pro Gln Leu Ser Leu Asp Ala Gln Leu Pro Ser Gly Lys
930 935 940

Val Glu Val Ala Gly Ala Glu Ser Lys Pro Lys Gly Ser Arg Phe Ala
 945 950 955 960

Leu Pro Lys Phe Gly Ala Lys Gly Arg Asp Ser Glu Ala Asp Val Leu
 965 970 975

Val Ala Gly Glu Ala Glu Leu Glu Gly Lys Gly Trp Gly Trp Asp Gly
 980 985 990

Lys Val Lys Met Pro Lys Leu Lys Met Pro Ser Phe Gly Leu Ser Arg
 995 1000 1005

Gly Lys Glu Ala Glu Ile Gln Asp Gly Arg Val Ser Pro Gly Glu
 1010 1015 1020

Lys Leu Glu Ala Ile Ala Gly Gln Leu Lys Ile Pro Glu Val Glu
 1025 1030 1035

Leu Val Thr Pro Gly Ala Gln Glu Thr Glu Lys Val Thr Ser Gly
 1040 1045 1050

Val Lys Pro Ser Gly Leu Gln Val Ser Thr Thr Arg Gln Val Val
 1055 1060 1065

Ala Glu Gly Gln Glu Gly Ala Gln Arg Val Ser Ser Leu Gly Ile
 1070 1075 1080

Ser Leu Pro Gln Val Glu Leu Ala Ser Phe Gly Glu Ala Gly Pro
 1085 1090 1095

Glu Ile Ala Ala Pro Ser Ala Glu Gly Thr Val Gly Ser Arg Ile
 1100 1105 1110

Gln Val Pro Gln Val Met Leu Glu Leu Pro Gly Thr Gln Val Ala
 1115 1120 1125

Gly Gly Asp Leu Leu Val Gly Glu Gly Ile Phe Lys Met Pro Thr
 1130 1135 1140

Val Thr Val Pro Gln Leu Glu Leu Asp Val Gly Leu Gly His Glu
 1145 1150 1155

Ala Gln Ala Gly Glu Thr Ala Lys Ser Glu Gly Gly Leu Lys Leu
 1160 1165 1170

Lys Leu Pro Thr Leu Gly Ala Gly Gly Lys Gly Glu Gly Ala Glu
1175 1180 1185

Ala Gln Ser Pro Glu Ala Gln His Thr Phe His Ile Ser Leu Pro
1190 1195 1200

Asp Val Glu Leu Thr Ser Pro Val Ser Ser His Ala Glu Tyr Gln
1205 1210 1215

Val Val Glu Gly Asp Gly Asp Gly Gly His Lys Leu Lys Val Arg
1220 1225 1230

Leu Pro Leu Phe Gly Leu Ala Arg Ala Lys Glu Gly Ile Glu Thr
1235 1240 1245

Gly Glu Lys Val Lys Ser Pro Lys Leu Arg Leu Pro Arg Val Gly
1250 1255 1260

Phe Ser Gln Ser Glu Ser Ala Ser Gly Glu Gly Ser Pro Ser Pro
1265 1270 1275

Glu Glu Glu Glu Glu Gly Ser Gly Glu Gly Ala Ser Gly Arg Arg
1280 1285 1290

Gly Arg Val Arg Val Arg Leu Pro Arg Val Gly Leu Ala Ser Pro
1295 1300 1305

Ser Lys Gly Ser Lys Gly Gln Glu Gly Asp Ala Ala Ser Lys Ser
1310 1315 1320

Pro Val Gly Glu Lys Ser Pro Lys Phe Arg Phe Pro Arg Val Ser
1325 1330 1335

Leu Ser Pro Lys Ala Arg Ser Gly Ser Lys Asp Arg Glu Glu Gly
1340 1345 1350

Gly Phe Arg Val Arg Leu Pro Ser Val Gly Phe Ser Glu Thr Ala
1355 1360 1365

Ala Pro Gly Ser Ala Arg Ile Glu Gly Thr Gln Ala Ala Ala Ile
1370 1375 1380

<210> 87
<211> 1383
<212> PRT
<213> Rat

<400> 87

Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
1 5 10 15

Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Phe
20 25 30

Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg
35 40 45

Glu Asp Ser Pro Ala Ala Lys Ser Leu Ser Leu Gln Glu Gly Asp Gln
50 55 60

Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala
65 70 75 80

Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu
85 90 95

Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val
100 105 110

Ser Gly Tyr Glu Met Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Asn
115 120 125

Ile Gln Ser Leu Ser Pro Val Lys Lys Lys Lys Met Val Ile Gly Thr
130 135 140

Leu Gly Thr Pro Ala Asp Leu Ala Pro Val Asp Val Glu Phe Ser Phe
145 150 155 160

Pro Lys Phe Ser Arg Leu Arg Arg Gly Leu Lys Ala Asp Ala Val Lys
165 170 175

Gly Pro Val Pro Ala Ala Pro Ala Arg Arg Arg Leu Gln Leu Pro Arg
180 185 190

Leu Arg Val Arg Glu Val Ala Glu Glu Ala Gln Val Ala Arg Met Ala
195 200 205

Ala Ala Ala Pro Pro Ser Arg Lys Ala Lys Ser Glu Ala Glu Val Ala
210 215 220

Thr Gly Ala Gly Phe Thr Ala Pro Gln Ile Glu Leu Val Gly Pro Arg
225 230 235 240

Leu Pro Ser Ala Glu Val Gly Val Pro Lys Val Ser Val Pro Lys Gly
 245 250 255

Thr Pro Ser Thr Glu Ala Ala Ser Gly Phe Ala Leu His Leu Pro Thr
 260 265 270

Leu Gly Leu Gly Ala Pro Ala Ala Pro Ala Val Glu Pro Pro Thr Thr
 275 280 285

Gly Ile Gln Val Pro Gln Val Glu Leu Pro Thr Leu Pro Ser Leu Pro
 290 295 300

Thr Leu Pro Thr Leu Pro Cys Leu Asp Thr Gln Glu Gly Ala Ala Val
 305 310 315 320

Val Lys Val Pro Thr Leu Asp Val Ala Ala Pro Ser Val Glu Val Asp
 325 330 335

Leu Ala Leu Pro Gly Ala Glu Val Glu Ala Gln Gly Glu Val Pro Glu
 340 345 350

Val Ala Leu Lys Met Pro Arg Leu Ser Phe Pro Arg Phe Gly Val Arg
 355 360 365

Gly Lys Glu Ala Thr Glu Ala Lys Val Val Lys Gly Ser Pro Glu Ala
 370 375 380

Lys Ala Lys Gly Pro Arg Leu Arg Met Pro Thr Phe Gly Leu Ser Leu
 385 390 395 400

Leu Glu Ser Arg Pro Ser Gly Pro Glu Val Ala Ala Glu Ser Lys Leu
 405 410 415

Lys Leu Pro Thr Leu Lys Met Pro Ser Phe Gly Ile Ser Val Ala Gly
 420 425 430

Pro Glu Val Lys Ala Pro Lys Gly Pro Glu Val Lys Leu Pro Lys Val
 435 440 445

Pro Glu Ile Lys Leu Pro Lys Ala Pro Glu Ala Ala Ile Pro Asp Val
 450 455 460

Gln Leu Pro Glu Val Gln Leu Pro Lys Met Ser Asp Met Lys Leu Pro
 465 470 475 480

Lys Ile Pro Glu Met Ala Val Pro Asp Val His Leu Pro Glu Val Lys
485 490 495

Leu Pro Lys Val Pro Glu Met Lys Val Pro Glu Met Lys Leu Pro Lys
500 505 510

Ile Pro Glu Met Ala Val Pro Asp Val His Leu Pro Asp Ile Gln Leu
515 520 525

Pro Lys Val Pro Glu Met Lys Leu Pro Asp Met Lys Leu Pro Lys Val
530 535 540

Pro Glu Met Ala Val Pro Asp Val His Leu Pro Asp Ile Gln Leu Pro
545 550 555 560

Lys Val Pro Glu Met Lys Leu Pro Asp Met Lys Leu Pro Lys Val Pro
565 570 575

Glu Met Ala Val Pro Asp Val Arg Ile Pro Glu Val Gln Leu Pro Lys
580 585 590

Val Ser Glu Val Lys Leu Pro Lys Ile Pro Asp Met Ala Val Pro Asp
595 600 605

Val Arg Leu Pro Glu Leu Gln Leu Pro Lys Met Ser Glu Val Lys Leu
610 615 620

Pro Lys Ile Pro Asp Met Ala Val Pro Asp Val Arg Leu Pro Glu Val
625 630 635 640

Gln Leu Pro Lys Val Ser Glu Leu Lys Leu Pro Lys Val Pro Glu Met
645 650 655

Thr Met Pro Asp Ile Arg Leu Pro Glu Val Gln Leu Pro Lys Val Pro
660 665 670

Asp Ile Lys Leu Pro Glu Ile Lys Leu Pro Lys Val Pro Glu Met Ala
675 680 685

Val Pro Asp Val Pro Leu Pro Glu Leu Gln Leu Pro Lys Val Pro Gln
690 695 700

Val Pro Asp Val His Leu Pro Lys Val Pro Glu Met Lys Leu Pro Lys
705 710 715 720

Val Pro Glu Ala Gln Arg Lys Ser Ala Gly Ala Glu Gln Ala Glu Lys

| | | |
|--|-----|-----|
| 725 | 730 | 735 |
| Thr Glu Phe Ser Phe Lys Leu Pro Lys Met Thr Val Pro Lys Leu Gly 740 745 750 | | |
| Lys Val Thr Lys Pro Gly Glu Ala Gly Ile Glu Val Pro Asp Lys Leu 755 760 765 | | |
| Leu Ile Leu Pro Cys Leu Gln Pro Glu Val Gly Thr Glu Val Ala Arg 770 775 780 | | |
| Val Gly Val Pro Ser Leu Ser Leu Pro Ser Val Glu Leu Asp Leu Pro 785 790 795 800 | | |
| Gly Ala Leu Gly Leu Glu Gly Gln Val Gln Glu Ala Val Ser Gly Lys 805 810 815 | | |
| Val Glu Lys Pro Glu Gly Pro Arg Val Ala Val Gly Thr Gly Glu Ala 820 825 830 | | |
| Gly Phe Arg Val Pro Ser Val Glu Ile Val Asn Pro Gln Leu Pro Thr 835 840 845 | | |
| Val Glu Val Lys Lys Glu Gln Leu Glu Met Val Glu Met Lys Val Lys 850 855 860 | | |
| Pro Thr Ser Lys Phe Ser Leu Pro Lys Phe Gly Leu Ser Gly Pro Lys 865 870 875 880 | | |
| Ala Val Lys Ala Glu Val Glu Gly Pro Gly Arg Ala Thr Lys Leu Lys 885 890 895 | | |
| Val Ser Lys Phe Ala Ile Ser Leu Pro Arg Ala Arg Ala Gly Thr Asp 900 905 910 | | |
| Ala Asp Ala Lys Gly Ala Gly Glu Ala Gly Leu Leu Pro Ala Leu Asp 915 920 925 | | |
| Leu Ser Ile Pro Gln Leu Ser Leu Asp Ala Gln Leu Pro Ser Gly Lys 930 935 940 | | |
| Val Glu Val Ala Gly Ala Glu Ser Lys Pro Lys Gly Ser Arg Phe Ala 945 950 955 960 | | |
| Leu Pro Lys Phe Gly Ala Lys Gly Arg Asp Ser Glu Ala Asp Val Leu 965 970 975 | | |

Val Ala Gly Glu Ala Glu Leu Glu Gly Lys Gly Trp Gly Trp Asp Gly
980 985 990

Lys Val Lys Met Pro Lys Leu Lys Met Pro Ser Phe Gly Leu Ser Arg
995 1000 1005

Gly Lys Glu Ala Glu Ile Gln Asp Gly Arg Val Ser Pro Gly Glu
1010 1015 1020

Lys Leu Glu Ala Ile Ala Gly Gln Leu Lys Ile Pro Glu Val Glu
1025 1030 1035

Leu Val Thr Pro Gly Ala Gln Glu Thr Glu Lys Val Thr Ser Gly
1040 1045 1050

Val Lys Pro Ser Gly Leu Gln Val Ser Thr Thr Arg Gln Val Val
1055 1060 1065

Ala Glu Gly Gln Glu Gly Ala Gln Arg Val Ser Ser Leu Gly Ile
1070 1075 1080

Ser Leu Pro Gln Val Glu Leu Ala Ser Phe Gly Glu Ala Gly Pro
1085 1090 1095

Glu Ile Ala Ala Pro Ser Ala Glu Gly Thr Val Gly Ser Arg Ile
1100 1105 1110

Gln Val Pro Gln Val Met Leu Glu Leu Pro Gly Thr Gln Val Ala
1115 1120 1125

Gly Gly Asp Leu Leu Val Gly Glu Gly Ile Phe Lys Met Pro Thr
1130 1135 1140

Val Thr Val Pro Gln Leu Glu Leu Asp Val Gly Leu Gly His Glu
1145 1150 1155

Ala Gln Ala Gly Glu Thr Ala Lys Ser Glu Gly Gly Leu Lys Leu
1160 1165 1170

Lys Leu Pro Thr Leu Gly Ala Gly Gly Lys Gly Glu Gly Ala Glu
1175 1180 1185

Ala Gln Ser Pro Glu Ala Gln His Thr Phe His Ile Ser Leu Pro
1190 1195 1200

Asp Val Glu Leu Thr Ser Pro Val Ser Ser His Ala Glu Tyr Gln
 1205 1210 1215

Val Val Glu Gly Asp Gly Asp Gly Gly His Lys Leu Lys Val Arg
 1220 1225 1230

Leu Pro Leu Phe Gly Leu Ala Arg Ala Lys Glu Gly Ile Glu Thr
 1235 1240 1245

Gly Glu Lys Val Lys Ser Pro Lys Leu Arg Leu Pro Arg Val Gly
 1250 1255 1260

Phe Ser Gln Ser Glu Ser Ala Ser Gly Glu Gly Ser Pro Ser Pro
 1265 1270 1275

Glu Glu Glu Glu Glu Gly Ser Gly Glu Gly Ala Ser Gly Arg Arg
 1280 1285 1290

Gly Arg Val Arg Val Arg Leu Pro Arg Val Gly Leu Ala Ser Pro
 1295 1300 1305

Ser Lys Gly Ser Lys Gly Gln Glu Gly Asp Ala Ala Ser Lys Ser
 1310 1315 1320

Pro Val Gly Glu Lys Ser Pro Lys Phe Arg Phe Pro Arg Val Ser
 1325 1330 1335

Leu Ser Pro Lys Ala Arg Ser Gly Ser Lys Asp Arg Glu Glu Gly
 1340 1345 1350

Gly Phe Arg Val Arg Leu Pro Ser Val Gly Phe Ser Glu Thr Ala
 1355 1360 1365

Ala Pro Gly Ser Ala Arg Ile Glu Gly Thr Gln Ala Ala Ala Ile
 1370 1375 1380

<210> 88
 <211> 1461
 <212> PRT
 <213> Human

<400> 88

Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
 1 5 10 15

Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Ile

| | | |
|--|----|----|
| 20 | 25 | 30 |
| Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg 35 40 45 | | |
| Glu Asp Ser Pro Ala Ala Arg Ser Leu Ser Leu Gln Glu Gly Asp Gln 50 55 60 | | |
| Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala 65 70 75 80 | | |
| Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu 85 90 95 | | |
| Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val 100 105 110 | | |
| Ser Gly Tyr Glu Ile Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Asn 115 120 125 | | |
| Ile Gln Ser Leu Ser Pro Val Lys Lys Lys Lys Met Val Pro Gly Ala 130 135 140 | | |
| Leu Gly Val Pro Ala Asp Leu Ala Pro Val Asp Val Glu Phe Ser Phe 145 150 155 160 | | |
| Pro Lys Phe Ser Arg Leu Arg Arg Gly Leu Lys Ala Glu Ala Val Lys 165 170 175 | | |
| Gly Pro Val Pro Ala Ala Pro Ala Arg Arg Arg Leu Gln Leu Pro Arg 180 185 190 | | |
| Leu Arg Val Arg Glu Val Ala Glu Glu Ala Gln Ala Ala Arg Leu Ala 195 200 205 | | |
| Ala Ala Ala Pro Pro Pro Arg Lys Ala Lys Val Glu Ala Glu Val Ala 210 215 220 | | |
| Ala Gly Ala Arg Phe Thr Ala Pro Gln Val Glu Leu Val Gly Pro Arg 225 230 235 240 | | |
| Leu Pro Gly Ala Glu Val Gly Val Pro Gln Val Ser Ala Pro Lys Ala 245 250 255 | | |
| Ala Pro Ser Ala Glu Ala Ala Gly Gly Phe Ala Leu His Leu Pro Thr 260 265 270 | | |

Leu Gly Leu Gly Ala Pro Ala Pro Pro Ala Val Glu Ala Pro Ala Val
275 280 285

Gly Ile Gln Val Pro Gln Val Glu Leu Pro Ala Leu Pro Ser Leu Pro
290 295 300

Thr Leu Pro Thr Leu Pro Cys Leu Glu Thr Arg Glu Gly Ala Val Ser
305 310 315 320

Val Val Val Pro Thr Leu Asp Val Ala Ala Pro Thr Val Gly Val Asp
325 330 335

Leu Ala Leu Pro Gly Ala Glu Val Glu Ala Arg Gly Glu Ala Pro Glu
340 345 350

Val Ala Leu Lys Met Pro Arg Leu Ser Phe Pro Arg Phe Gly Ala Arg
355 360 365

Ala Lys Glu Val Ala Glu Ala Lys Val Ala Lys Val Ser Pro Glu Ala
370 375 380

Arg Val Lys Gly Pro Arg Leu Arg Met Pro Thr Phe Gly Leu Ser Leu
385 390 395 400

Leu Glu Pro Arg Pro Ala Ala Pro Glu Val Val Glu Ser Lys Leu Lys
405 410 415

Leu Pro Thr Ile Lys Met Pro Ser Leu Gly Ile Gly Val Ser Gly Pro
420 425 430

Glu Val Lys Val Pro Lys Gly Pro Glu Val Lys Leu Pro Lys Ala Pro
435 440 445

Glu Val Lys Leu Pro Lys Val Pro Glu Ala Ala Leu Pro Glu Val Arg
450 455 460

Leu Pro Glu Val Glu Leu Pro Lys Val Ser Glu Met Lys Leu Pro Lys
465 470 475 480

Val Pro Glu Met Ala Val Pro Glu Val Arg Leu Pro Glu Val Glu Leu
485 490 495

Pro Lys Val Ser Glu Met Lys Leu Pro Lys Val Pro Glu Met Ala Val
500 505 510

Pro Glu Val Arg Leu Pro Glu Val Gln Leu Leu Lys Val Ser Glu Met
515 520 525

Lys Leu Pro Lys Val Pro Glu Met Ala Val Pro Glu Val Arg Leu Pro
530 535 540

Glu Val Gln Leu Pro Lys Val Ser Glu Met Lys Leu Pro Glu Val Ser
545 550 555 560

Glu Val Ala Val Pro Glu Val Arg Leu Pro Glu Val Gln Leu Pro Lys
565 570 575

Val Pro Glu Met Lys Val Pro Glu Met Lys Leu Pro Lys Val Pro Glu
580 585 590

Met Lys Leu Pro Glu Met Lys Leu Pro Glu Val Gln Leu Pro Lys Val
595 600 605

Pro Glu Met Ala Val Pro Asp Val His Leu Pro Glu Val Gln Leu Pro
610 615 620

Lys Val Pro Glu Met Lys Leu Pro Glu Met Lys Leu Pro Glu Val Lys
625 630 635 640

Leu Pro Lys Val Pro Glu Met Ala Val Pro Asp Val His Leu Pro Glu
645 650 655

Val Gln Leu Pro Lys Val Pro Glu Met Lys Leu Pro Lys Met Pro Glu
660 665 670

Met Ala Val Pro Glu Val Arg Leu Pro Glu Val Gln Leu Pro Lys Val
675 680 685

Ser Glu Met Lys Leu Pro Lys Val Pro Glu Met Ala Val Pro Asp Val
690 695 700

His Leu Pro Glu Val Gln Leu Pro Lys Val Cys Glu Met Lys Val Pro
705 710 715 720

Asp Met Lys Leu Pro Glu Ile Lys Leu Pro Lys Val Pro Glu Met Ala
725 730 735

Val Pro Asp Val His Leu Pro Glu Val Gln Leu Pro Lys Val Ser Glu
740 745 750

Ile Arg Leu Pro Glu Met Gln Val Pro Lys Val Pro Asp Val His Leu
 755 760 765
 Pro Lys Ala Pro Glu Val Lys Leu Pro Arg Ala Pro Glu Val Gln Leu
 770 775 780
 Lys Ala Thr Lys Ala Glu Gln Ala Glu Gly Met Glu Phe Gly Phe Lys
 785 790 795 800
 Met Pro Lys Met Thr Met Pro Lys Leu Gly Arg Ala Glu Ser Pro Ser
 805 810 815
 Arg Gly Lys Pro Gly Glu Ala Gly Ala Glu Val Ser Gly Lys Leu Val
 820 825 830
 Thr Leu Pro Cys Leu Gln Pro Glu Val Asp Gly Glu Ala His Val Gly
 835 840 845
 Val Pro Ser Leu Thr Leu Pro Ser Val Glu Leu Asp Leu Pro Gly Ala
 850 855 860
 Leu Gly Leu Gln Gly Gln Val Pro Ala Ala Lys Met Gly Lys Gly Glu
 865 870 875 880
 Arg Val Glu Gly Pro Glu Val Ala Ala Gly Val Arg Glu Val Gly Phe
 885 890 895
 Arg Val Pro Ser Val Glu Ile Val Thr Pro Gln Leu Pro Ala Val Glu
 900 905 910
 Ile Glu Glu Gly Arg Leu Glu Met Ile Glu Thr Lys Val Lys Pro Ser
 915 920 925
 Ser Lys Phe Ser Leu Pro Lys Phe Gly Leu Ser Gly Pro Lys Val Ala
 930 935 940
 Lys Ala Glu Ala Glu Gly Ala Gly Arg Ala Thr Lys Leu Lys Val Ser
 945 950 955 960
 Lys Phe Ala Ile Ser Leu Pro Lys Ala Arg Val Gly Ala Glu Ala Glu
 965 970 975
 Ala Lys Gly Ala Gly Glu Ala Gly Leu Leu Pro Ala Leu Asp Leu Ser
 980 985 990
 Ile Pro Gln Leu Ser Leu Asp Ala His Leu Pro Ser Gly Lys Val Glu

| 995 | 1000 | 1005 |
|---------------------------------|-----------------------------|-------------------------|
| Val Ala Gly Ala Asp Leu 1010 | Lys Phe Lys Gly Pro 1015 | Arg Phe Ala Leu 1020 |
| Pro Lys Phe Gly Val Arg 1025 | Gly Arg Asp Thr Glu 1030 | Ala Ala Glu Leu 1035 |
| Val Pro Gly Val Ala Glu 1040 | Leu Glu Gly Lys Gly 1045 | Trp Gly Trp Asp 1050 |
| Gly Arg Val Lys Met Pro 1055 | Lys Leu Lys Met Pro 1060 | Ser Phe Gly Leu 1065 |
| Ala Arg Gly Lys Glu Ala 1070 | Glu Val Gln Gly Asp 1075 | Arg Ala Ser Pro 1080 |
| Gly Glu Lys Ala Glu Ser 1085 | Thr Ala Val Gln Leu 1090 | Lys Ile Pro Glu 1095 |
| Val Glu Leu Val Thr Leu 1100 | Gly Ala Gln Glu Glu 1105 | Gly Arg Ala Glu 1110 |
| Gly Ala Val Ala Val Ser 1115 | Gly Met Gln Leu Ser 1120 | Gly Leu Lys Val 1125 |
| Ser Thr Ala Arg Gln Val 1130 | Val Thr Glu Gly His 1135 | Asp Ala Gly Leu 1140 |
| Arg Met Pro Pro Leu Gly 1145 | Ile Ser Leu Pro Gln 1150 | Val Glu Leu Thr 1155 |
| Gly Phe Gly Glu Ala Gly 1160 | Thr Pro Gly Gln Gln 1165 | Ala Gln Ser Thr 1170 |
| Val Pro Ser Ala Glu Gly 1175 | Thr Ala Gly Tyr Arg 1180 | Val Gln Val Pro 1185 |
| Gln Val Thr Leu Ser Leu 1190 | Pro Gly Ala Gln Val 1195 | Ala Gly Gly Glu 1200 |
| Leu Leu Val Gly Glu Gly 1205 | Val Phe Lys Met Pro 1210 | Thr Val Thr Val 1215 |
| Pro Gln Leu Glu Leu Asp 1220 | Val Gly Leu Ser Arg 1225 | Glu Ala Gln Ala 1230 |

| | | | | | | | | | | | | | | |
|------|-----|-----|-----|-----|-----|------|-----|-----|-----|-----|------|-----|-----|-----|
| Gly | Glu | Ala | Ala | Thr | Gly | Glu | Gly | Gly | Leu | Arg | Leu | Lys | Leu | Pro |
| 1235 | | | | | | 1240 | | | | | 1245 | | | |
| Thr | Leu | Gly | Ala | Arg | Ala | Arg | Val | Gly | Gly | Glu | Gly | Ala | Glu | Glu |
| 1250 | | | | | | 1255 | | | | | 1260 | | | |
| Gln | Pro | Pro | Gly | Ala | Glu | Arg | Thr | Phe | Cys | Leu | Ser | Leu | Pro | Asp |
| 1265 | | | | | | 1270 | | | | | 1275 | | | |
| Val | Glu | Leu | Ser | Pro | Ser | Gly | Gly | Asn | His | Ala | Glu | Tyr | Gln | Val |
| 1280 | | | | | | 1285 | | | | | 1290 | | | |
| Ala | Glu | Gly | Glu | Gly | Glu | Ala | Gly | His | Lys | Leu | Lys | Val | Arg | Leu |
| 1295 | | | | | | 1300 | | | | | 1305 | | | |
| Pro | Arg | Phe | Gly | Leu | Val | Arg | Ala | Lys | Glu | Gly | Ala | Glu | Glu | Gly |
| 1310 | | | | | | 1315 | | | | | 1320 | | | |
| Glu | Lys | Ala | Lys | Ser | Pro | Lys | Leu | Arg | Leu | Pro | Arg | Val | Gly | Phe |
| 1325 | | | | | | 1330 | | | | | 1335 | | | |
| Ser | Gln | Ser | Glu | Met | Val | Thr | Gly | Glu | Gly | Ser | Pro | Ser | Pro | Glu |
| 1340 | | | | | | 1345 | | | | | 1350 | | | |
| Glu | Glu | Glu | Glu | Glu | Glu | Glu | Glu | Gly | Ser | Gly | Glu | Gly | Ala | Ser |
| 1355 | | | | | | 1360 | | | | | 1365 | | | |
| Gly | Arg | Arg | Gly | Arg | Val | Arg | Val | Arg | Leu | Pro | Arg | Val | Gly | Leu |
| 1370 | | | | | | 1375 | | | | | 1380 | | | |
| Ala | Ala | Pro | Ser | Lys | Ala | Ser | Arg | Gly | Gln | Glu | Gly | Asp | Ala | Ala |
| 1385 | | | | | | 1390 | | | | | 1395 | | | |
| Pro | Lys | Ser | Pro | Val | Arg | Glu | Lys | Ser | Pro | Lys | Phe | Arg | Phe | Pro |
| 1400 | | | | | | 1405 | | | | | 1410 | | | |
| Arg | Val | Ser | Leu | Ser | Pro | Lys | Ala | Arg | Ser | Gly | Ser | Gly | Asp | Gln |
| 1415 | | | | | | 1420 | | | | | 1425 | | | |
| Glu | Glu | Gly | Gly | Leu | Arg | Val | Arg | Leu | Pro | Ser | Val | Gly | Phe | Ser |
| 1430 | | | | | | 1435 | | | | | 1440 | | | |
| Glu | Thr | Gly | Ala | Pro | Gly | Pro | Ala | Arg | Met | Glu | Gly | Ala | Gln | Ala |
| 1445 | | | | | | 1450 | | | | | 1455 | | | |

Ala Ala Val
1460

<210> 89
<211> 147
<212> PRT
<213> Human

<400> 89

Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
1 5 10 15

Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Ile
20 25 30

Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg
35 40 45

Glu Asp Ser Pro Ala Ala Arg Ser Leu Ser Leu Gln Glu Gly Asp Gln
50 55 60

Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala
65 70 75 80

Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu
85 90 95

Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val
100 105 110

Ser Gly Tyr Glu Ile Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Val
115 120 125

Arg Val Leu Ser Pro Ala Pro Ala Leu Asp Cys Pro Ser Asp Pro Val
130 135 140

Ser Ala Pro
145

<210> 90
<211> 370
<212> PRT
<213> Arabidopsisthaliana

<400> 90

Met Ala Leu Met Lys Lys Ser Leu Ser Ala Ala Leu Leu Ser Ser Pro
1 5 10 15

Leu Leu Ile Ile Cys Leu Ile Ala Leu Leu Ala Asp Pro Phe Ser Val
 20 25 30

Gly Ala Arg Arg Leu Leu Glu Asp Pro Lys Pro Glu Ile Pro Lys Leu
 35 40 45

Pro Glu Leu Pro Lys Phe Glu Val Pro Lys Leu Pro Glu Phe Pro Lys
 50 55 60

Pro Glu Leu Pro Lys Leu Pro Glu Phe Pro Lys Pro Glu Leu Pro Lys
 65 70 75 80

Ile Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Val Pro Glu Ile Pro
 85 90 95

Lys Pro Glu Glu Thr Lys Leu Pro Asp Ile Pro Lys Leu Glu Leu Pro
 100 105 110

Lys Phe Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Met Pro Glu Ile
 115 120 125

Pro Lys Pro Glu Leu Pro Lys Val Pro Glu Ile Gln Lys Pro Glu Leu
 130 135 140

Pro Lys Met Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Phe Pro Glu
 145 150 155 160

Ile Pro Lys Pro Asp Leu Pro Lys Phe Pro Glu Asn Ser Lys Pro Glu
 165 170 175

Val Pro Lys Leu Met Glu Thr Glu Lys Pro Glu Ala Pro Lys Val Pro
 180 185 190

Glu Ile Pro Lys Pro Glu Leu Pro Lys Leu Pro Glu Val Pro Lys Leu
 195 200 205

Glu Ala Pro Lys Val Pro Glu Ile Gln Lys Pro Glu Leu Pro Lys Met
 210 215 220

Pro Glu Leu Pro Lys Met Pro Glu Ile Gln Lys Pro Glu Leu Pro Lys
 225 230 235 240

Leu Pro Glu Val Pro Lys Leu Glu Ala Pro Lys Val Pro Glu Ile Gln
 245 250 255

Lys Pro Glu Leu Pro Lys Met Pro Glu Leu Pro Lys Met Pro Glu Ile
 260 265 270

Gln Lys Pro Glu Leu Pro Lys Met Pro Glu Ile Gln Lys Pro Glu Leu
 275 280 285

Pro Lys Val Pro Glu Val Pro Lys Pro Glu Leu Pro Thr Val Pro Glu
 290 295 300

Val Pro Lys Ser Glu Ala Pro Lys Phe Pro Glu Ile Pro Lys Pro Glu
 305 310 315 320

Leu Pro Lys Ile Pro Glu Val Pro Lys Pro Glu Leu Pro Lys Val Pro
 325 330 335

Glu Ile Thr Lys Pro Ala Val Pro Glu Ile Pro Lys Pro Glu Leu Pro
 340 345 350

Thr Met Pro Gln Leu Pro Lys Leu Pro Glu Phe Pro Lys Val Pro Gly
 355 360 365

Thr Pro
 370

<210> 91
 <211> 370
 <212> PRT
 <213> Arabidopsisthaliana

<400> 91

Met Ala Leu Met Lys Lys Ser Leu Ser Ala Ala Leu Leu Ser Ser Pro
 1 5 10 15

Leu Leu Ile Ile Cys Leu Ile Ala Leu Leu Ala Asp Pro Phe Ser Val
 20 25 30

Gly Ala Arg Arg Leu Leu Glu Asp Pro Lys Pro Glu Ile Pro Lys Leu
 35 40 45

Pro Glu Leu Pro Lys Phe Glu Val Pro Lys Leu Pro Glu Phe Pro Lys
 50 55 60

Pro Glu Leu Pro Lys Leu Pro Glu Phe Pro Lys Pro Glu Leu Pro Lys
 65 70 75 80

Ile Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Val Pro Glu Ile Pro

85

90

95

Lys Pro Glu Glu Thr Lys Leu Pro Asp Ile Pro Lys Leu Glu Leu Pro
 100 105 110

Lys Phe Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Met Pro Glu Ile
 115 120 125

Pro Lys Pro Glu Leu Pro Lys Val Pro Glu Ile Gln Lys Pro Glu Leu
 130 135 140

Pro Lys Met Pro Glu Ile Pro Lys Pro Glu Leu Pro Lys Phe Pro Glu
 145 150 155 160

Ile Pro Lys Pro Asp Leu Pro Lys Phe Pro Glu Asn Ser Lys Ser Glu
 165 170 175

Val Pro Lys Leu Met Glu Thr Glu Lys Pro Glu Ala Pro Lys Val Pro
 180 185 190

Glu Ile Pro Lys Pro Glu Leu Pro Lys Leu Pro Glu Val Pro Lys Leu
 195 200 205

Glu Ala Pro Lys Val Pro Glu Ile Gln Lys Pro Glu Leu Pro Lys Met
 210 215 220

Pro Glu Leu Pro Lys Met Pro Glu Ile Gln Lys Pro Glu Leu Pro Lys
 225 230 235 240

Leu Pro Glu Val Pro Lys Leu Glu Ala Pro Lys Val Pro Glu Ile Gln
 245 250 255

Lys Pro Glu Leu Pro Lys Met Pro Glu Leu Pro Lys Met Pro Glu Ile
 260 265 270

Gln Lys Pro Glu Leu Pro Lys Met Pro Glu Ile Gln Lys Pro Glu Leu
 275 280 285

Pro Lys Val Pro Glu Val Pro Lys Pro Glu Leu Pro Thr Val Pro Glu
 290 295 300

Val Pro Lys Ser Glu Ala Pro Lys Phe Pro Glu Ile Pro Lys Pro Glu
 305 310 315 320

Leu Pro Lys Ile Pro Glu Val Pro Lys Pro Glu Leu Pro Lys Val Pro
 325 330 335

Glu Ile Thr Lys Pro Ala Val Pro Glu Ile Pro Lys Pro Glu Leu Pro
 340 345 350

Thr Met Pro Gln Leu Pro Lys Leu Pro Glu Phe Pro Lys Val Pro Gly
 355 360 365

Thr Pro
 370

<210> 92
 <211> 147
 <212> PRT
 <213> Human

<400> 92

Met Glu Ala Arg Ser Arg Ser Ala Glu Glu Leu Arg Arg Ala Glu Leu
 1 5 10 15

Val Glu Ile Ile Val Glu Thr Glu Ala Gln Thr Gly Val Ser Gly Ile
 20 25 30

Asn Val Ala Gly Gly Gly Lys Glu Gly Ile Phe Val Arg Glu Leu Arg
 35 40 45

Glu Asp Ser Pro Ala Ala Arg Ser Leu Ser Leu Gln Glu Gly Asp Gln
 50 55 60

Leu Leu Ser Ala Arg Val Phe Phe Glu Asn Phe Lys Tyr Glu Asp Ala
 65 70 75 80

Leu Arg Leu Leu Gln Cys Ala Glu Pro Tyr Lys Val Ser Phe Cys Leu
 85 90 95

Lys Arg Thr Val Pro Thr Gly Asp Leu Ala Leu Arg Pro Gly Thr Val
 100 105 110

Ser Gly Tyr Glu Ile Lys Gly Pro Arg Ala Lys Val Ala Lys Leu Val
 115 120 125

Arg Val Leu Ser Pro Ala Pro Ala Leu Asp Cys Pro Ser Asp Pro Val
 130 135 140

Ser Ala Pro
 145

<210> 93
 <211> 214
 <212> PRT
 <213> Human

<400> 93

Met Glu Leu Leu Gly Glu Gly Ala Ile Leu Gln Gly Arg Arg Glu Ser
 1 5 10 15

Gln Met Glu Ala Ala Pro Gly Ile Gln Thr Cys Gly His Ser Ala Glu
 20 25 30

Leu Pro Ser Gln Gly Met Gly Arg Thr Arg Ala Glu Arg Ala Thr Ser
 35 40 45

Pro Val Arg Pro Ser Ile Thr Trp Lys Ile Gly Ser Pro Lys Val Asp
 50 55 60

Gly Arg His Thr Pro Met Pro Phe Pro Ser Val Ser Thr Gly Glu Gly
 65 70 75 80

Lys Ser Thr Leu Trp Ile Leu Tyr Leu His Cys Phe Gly Ser Arg Lys
 85 90 95

Ser Pro Asp Phe Ser Thr Pro Pro Arg Glu Pro Lys Ser Gln Gly Met
 100 105 110

Leu Lys Glu Gln Ala Arg Lys Met Arg Gly Gln Arg Gly Gly Arg Glu
 115 120 125

Gly Ala Lys Gly Thr Leu Lys Thr Gln Arg Pro Pro Ser Lys Asp Gln
 130 135 140

Ala Pro Leu Ala His Gly Pro Arg Glu Lys Gln Val Pro Ala Asp Glu
 145 150 155 160

Ser Phe Leu Gln Lys Pro Arg Leu Pro Asp Leu Val Lys Gln Gln Pro
 165 170 175

Asn Arg Ser Leu Ser Thr Asn Val Arg Gly Ala Glu Pro Ser Pro Ser
 180 185 190

Leu Ala Thr Glu Leu Val Leu Lys Lys Leu Val Pro Ala Ser Thr Cys
 195 200 205

Gln Glu Leu Pro Lys Thr
 210